

Original Paper

## Beneficial Effects of Low Frequency Vibration on Human Chondrocytes *in Vitro*

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### Key Words

Vibration • Chondrocytes • Extracellular matrix • Apoptosis • Focal adhesion • Annexin A2

### Abstract

**Background/Aims:** In articular cartilage, chondrocytes are the predominant cell type. A long-term stay in space can lead to bone loss and cartilage breakdown. Due to the poor regenerative capacity of cartilage, this may impair the crewmembers' mobility and influence mission activities. Beside microgravity other factors such as cosmic radiation and vibration might be important for cartilage degeneration. Vibration at different frequencies showed various effects on cartilage *in vivo*, but knowledge about its impact on chondrocytes *in vitro* is sparse. **Methods:** Human chondrocytes were exposed to a vibration device, simulating the vibration profile occurring during parabolic flights, for 24 h (VIB) and compared to static controls. Phase-contrast microscopy, immunofluorescence, F-actin and TUNEL staining as well as quantitative real-time PCR were performed to examine effects on morphology, cell viability and shape as well as gene expression. The results were compared to earlier studies using semantic analyses. **Results:** No morphological changes or cytoskeletal alterations were observed in VIB and no apoptotic cells were found. A reorganization and increase in fibronectin were detected in VIB samples by immunofluorescence technique. *PXN*, *VCL*, *ANXA1*, *ANXA2*, *BAX*, and *BCL2* revealed differential regulations. **Conclusion:** Long-term VIB did not damage human chondrocytes *in vitro*. The reduction of *ANXA2*, and up-regulation of *ANXA1*, *PXN* and *VCL* mRNAs suggest that long-term vibration might even positively influence cultured chondrocytes.

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## Introduction

Cartilage cells, also called chondrocytes, are the sole cellular component in cartilage tissue. They synthesize extracellular matrix (ECM) proteins, such as collagen II, proteoglycans, fibronectin and others.

Several publications demonstrated that mechanical stress influences human chondrocytes *in vitro* and alters various biological processes such as proliferation, cell adhesion, differentiation, and signal transduction [1, 2]. Human chondrocytes exposed to parabolic flight (PF) maneuvers showed an up-regulation of the cytoskeletal network genes and proteins suggesting that short-term microgravity ( $\mu g$ ) influences the cells to respond by reorganizing the cytoskeleton [3]. Similar data were found when human chondrocytes were investigated for 24 h on the random positioning machine (RPM). The chondrocytes had changed their ECM synthesis while they reorganized their cytoskeleton prior to forming three-dimensional (3D) aggregates [4, 5].

In earlier studies an effect of vibration (VIB) was observed, when chondrocytes of human origin were investigated for 2 h and 24 h with vibraplex device developed by the German Aerospace Center (Deutsches Zentrum für Luft- und Raumfahrt, DLR) [3, 6]. Short-term VIB did not negatively influence the cartilage cells [3] and also long-term VIB did not alter the morphology of human chondrocytes *in vitro*. No signs of apoptosis were found. However, a reduction of osteopontin protein and a decrease in *PSMD4* and *TBX15* gene expression suggested that VIB might have positive effects on human chondrocytes [6].

Whole-body vibration (WBV) is currently under investigation to examine whether it is an option to treat the musculoskeletal disease osteoarthritis (OA). The effects are not clarified. A recent study showed that WBV induced cartilage degeneration in mice [7], while the application of lower frequencies of WBV showed a more differentiated picture. WBV of 10 Hz and 20 Hz resulted in a reduced cartilage resorption, an increased cartilage formation, and a delayed cartilage degradation, especially when applying 20 Hz, whereas higher frequencies of 30 Hz and 40 Hz induced worsening limb function, shrinking cartilage volume and cartilage resorption [8].

The principal aim of this study was first to address changes in the expression of selected genes of the ECM, focal adhesions and pathways of apoptosis in human chondrocytes exposed to VIB using a Vibraplex device, and secondly to compare these data with earlier  $\mu g$  results from studies reporting about the effects of simulated microgravity (*s- $\mu g$* ) using a Random Positioning Machine (RPM) or of real microgravity (*r- $\mu g$* ) achieved during PF or space missions. For this purpose, *in silico* analyses were performed. Moreover, the cells were examined by immunofluorescence staining and laser scanning microscopy to visualize the F-actin cytoskeleton, as well as vimentin, fibronectin and ICAM-1 proteins.

## Materials and Methods

### *Cells and Culture Medium*

Human chondrocytes from six different donors were bought from the company Provitro (Berlin, Germany). The cells were cultured in Chondrocyte Growth Medium (CGM; Provitro®, Berlin, Germany) supplemented with 10 % fetal calf serum (Provitro®, Berlin, Germany) and antibiotics – 100 IU penicillin/mL and 100  $\mu g$  streptomycin/mL (Provitro®, Berlin, Germany). Chondrocytes of passage 3 grown in T25 cm<sup>2</sup> flasks (25 cm<sup>2</sup>; Sarstedt, Nümbrecht, Germany) were used for the experiments. The procedure was described in detail in [6].

For immunohistochemical studies, the cells were seeded in slide flasks (Thermo Fisher Scientific, Waltham, MA, USA) 24 h before the experiments (n=40 each group).

### *Vibration experiments*

The vibration experiments were conducted as described in [3]. Briefly, the Vibraplex is a VIB platform designed and constructed by the Deutsches Zentrum für Luft- und Raumfahrt (DLR, German Center for

Aerospace Medicine) - Cologne, and can be used for studies to mimic the mechanical VIB encountered on an airplane during PFs.

The Vibraplex VIB platform (frequency range 0.2 Hz-14 kHz) was used to create VIB like to the ones occurring during PFs. The corresponding vibrations were recorded and analyzed by Schmidt [9]. The device is driven by an amplified wave signal equal to a  $1/f^2$  noise (red noise), which was generated with the software WaveLab 4.0 from Steinberg.

The vibration treatment of the samples during the simulated free fall phase was alleviated by about -6 dB. This is half of the energy found during the simulated horizontal flight. During the simulated pull-up and pull-out phases, the vibrations were amplified by about +6 dB. The intensity of the noise applied was equivalent to  $100 \text{ W/m}^2$ . The Vibraplex uses 2W with an experimental area of about  $0.02 \text{ m}^2$ .

**Table 1.** Primary antibodies for immunofluorescence staining

Antibody	Dilution	Company
Vimentin Mouse	1:200	Sigma-Aldrich V5255
ICAM-1 Rabbit	1:25	Cell Signaling 4915
Fibronectin Rabbit	1:400	Sigma-Aldrich F3648

### *Immunofluorescence and F-actin staining*

After the experimental procedures (VIB and static control experiments), the cell culture medium was immediately aspirated and the chondrocytes were fixed with 4% PFA for 30 minutes at room temperature (RT). Then the slides were washed two times in DPBS. Afterwards, the slides were washed with 0.1% Triton X in DPBS for 10 min with agitation, which was followed by three more washing steps with DPBS. To prevent non-specific binding, the slides were incubated in 3% BSA in a wet chamber for one hour. Primary antibodies diluted in 1% BSA were added to the slides overnight at  $4^\circ \text{C}$  (Table 1). The following day, the slides were washed 2 times in DPBS and the secondary antibody Alexa Fluor plus 488 goat anti-mouse IgG (H + L) (1:400) or Alexa Fluor 488 F(ab')<sub>2</sub> fragment of goat anti-rabbit IgG (H + L) (1:500) (both Invitrogen by Thermo Fischer Scientific) diluted in 1% BSA was added for one hour at RT. Afterwards, the slides were washed again with DPBS and Alexa Fluor 568 phalloidin (Invitrogen by Thermo Fischer Scientific) was added for one hour, followed by three times washing with DPBS and mounting with Fluoroshield with DAPI (Sigma Life Science). Then the slides were incubated overnight at  $4^\circ \text{C}$ . A Carl ZEISS LSM 800 Confocal laser scanning microscope was used to examine the cells. Three lasers were used to examine the slides: 488 nm, 561 nm and 405 nm for visualization of Alexa 488, Alexa 568 and DAPI, respectively.

### *Viability staining*

TdT-mediated dUTP-biotin nick end labeling (TUNEL) staining was done according to the manual provided by the manufacturer (Thermo Fisher Scientific, Click-iT TUNEL Alexa Fluor 488 (cat# C10245)). The cells for 4',6-diamidino-2-phenylindole (DAPI) staining were fixed with 3.7% formaldehyde (room temperature, 10 min) and incubated in  $1 \mu\text{g/mL}$  DAPI in PBS for 15 min (Invitrogen/Molecular Probes, Darmstadt, Germany). Stained cell samples (VIB and static CON) were investigated utilizing a Leica DM 2000 microscope equipped with an objective with a calibrated magnification of 400x and connected to an external light source, Leica EL 6000 (Leica Microsystems GmbH, Wetzlar, Germany). To obtain positive controls the cells were treated with DNAase before the TUNEL staining. The results were shown in the inserts of Fig. 1C, D.

### *Microscopy*

The chondrocytes were examined with a Zeiss 510 META inverted confocal laser scanning microscope (Zeiss, Oberkochen, Germany), equipped with a Plan-Apochromat 363 1.4 objective [4]. Excitation and emission wavelengths were  $\lambda_{exc} = 488 \text{ nm}$  and  $\lambda_{em} = 505 \text{ nm}$  for FITC.

All samples were investigated with the image analysis program Scion Image (Version 1.63 Mac OS; Scion Corporation, Frederick, MD, USA). Phase contrast microscopy was performed using a LEICA DM2000 microscope equipped with a Leica DFC310 FX digital CCD color camera.

### *RNA isolation and quantitative real-time PCR (qPCR)*

The RNA isolation and qPCR were performed as published earlier [6, 10]. Briefly, RNA was isolated with the RNeasy Mini Kit (Qiagen, Hilden, Germany), with an additional DNase digestion step (Qiagen) in order to eliminate residual DNA contaminations. Subsequently, the amount of RNA was quantified via a Photometer Ultraspec2010 (Amersham Biosciences, Freiburg, Germany). The first strand cDNA synthesis

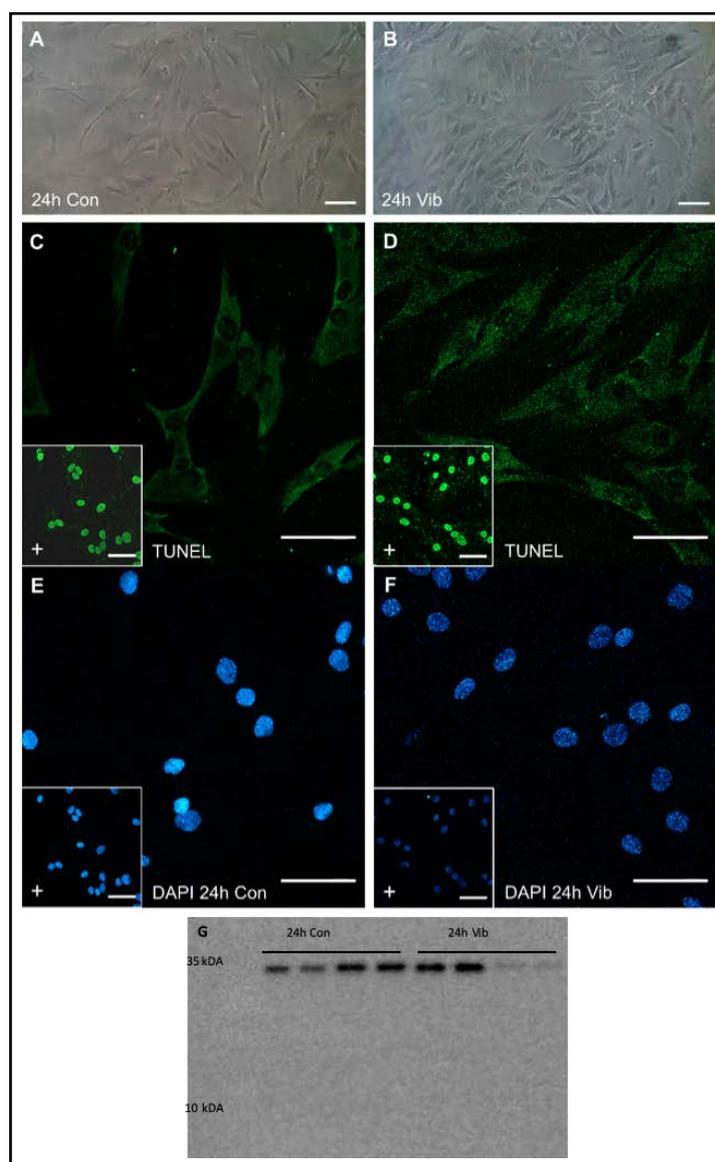
kit (Thermo Fisher Scientific, Waltham, US) was used for reverse transcription. qPCR was performed using the 7500 Fast Real-Time PCR System (Applied Biosystems, Darmstadt, Germany) according to routine protocols [6, 11]. cDNA-selective-primers were synthesized by TIB Molbiol (Berlin, Germany) and were listed in Table 2. The primers were designed using Primer Express (Applied Biosystems, Darmstadt, Germany) to have a  $T_m$  of  $\sim 60^\circ\text{C}$  and to span exon-exon boundaries. All samples were measured in triplicate. For normalization, 18S rRNA was used as a housekeeping gene. The comparative  $C_t$  ( $\Delta\Delta C_t$ ) method was used for relative quantification of transcription levels and non-vibrated controls were defined as 100 % for reference.

### Western blot analyses

Whole cell lysates were used for Western blotting following conventional protocols for gel electrophoresis and trans-blotting, as described earlier in [12]. Equal amounts of 10  $\mu\text{L}$  lysate containing 2  $\mu\text{g}/\mu\text{L}$  protein were loaded on precast TGX stain-free gels (Bio-Rad, Munich, Germany). Transturbo blot PVDF membranes (Bio-Rad) were used for blotting. An overview of the used antibodies and their applied concentrations is given in Table 3. The analysis was performed using ImageQuant™ LAS 4000 (GE Healthcare UK Limited, Buckinghamshire, UK), and the densitometry was performed using ImageJ (NIH).

### Pathway analyses

To study the mutual regulation of genes and to visualize localization and interactions between proteins, we entered relevant UniProtKB entry numbers in the Pathway Studio v.11 software (Elsevier Research Solutions, Amsterdam, The Netherlands). The graphs were generated for the gene expression and protein regulation and binding. The method was described in earlier studies [13, 14]. STRING analyses were done with STRING 11.0, available at <https://string-db.org/>.



**Fig. 1.** Morphology of human primary chondrocytes in (A) static controls and (B) after 24h of VIB. TUNEL assay in comparison to DAPI counter staining (C, D) was used for apoptosis detection (green fluorescence). Positive controls (inserts) were initiated by DNase 1 prior to staining. DAPI staining (E, F) after fixation with PFA indicated no apoptotic bodies inside the nuclei or decomposition. (G) No cleaved caspase-3 (17 kDa) could be detected by Western blot. The uncleaved Cas-3 band visible at 32 kDa. Scale bars: 100  $\mu\text{m}$ .

*Statistical analysis*

All statistical analyses were performed using SPSS 24 software (SPSS, Inc., Chicago, IL, USA). We used the Mann-Whitney-U-test to assess differences, which were considered significant at the level of  $p < 0.05$ . All data are presented as means  $\pm$  standard deviation.

**Results**

*Morphology of the chondrocytes*

After 24 h of VIB, no morphological differences between static control cells and vibrated chondrocytes were visible on commercially available human chondrocytes of passage 3 used for the experiments. The cells showed regular cell morphologies without any detectable alterations (Fig. 1A, B). Furthermore, all vibrated chondrocytes did not exhibit an increase in apoptosis (Fig. 1C-F). DAPI staining revealed normal nuclei in all samples (Fig. 1E, F). Positive controls given in the inserts showed apoptotic nuclei. In addition, Western blotting of cleaved caspase-3 demonstrated no activated caspase-3 (17 kDa) in all samples (Fig. 1G).

*The F-actin cytoskeleton, vimentin, fibronectin and ICAM-1*

Rhodamine-phalloidin staining was used to visualize the F-actin cytoskeleton after 24 h. Both static control as well as vibrated cells displayed a normal distribution and structure of actin fibers with no signs of damage, no disarrangements or deposits of F-actin at the cellular membranes (Fig. 2A, D, G, J, M, P).

The distribution and localization of the intermediate filament vimentin was similar in both groups (Fig. 2B, E).

Significant changes in the fibronectin content of chondrocytes exposed to mechanical vibration are shown in the Fig. 2H, K. A 24-hour-exposure to vibration showed that the cells started to secrete fibronectin into the extracellular space between adjacent cells, as can clearly be seen in Fig. 2K, L. An accumulation of this ECM protein was detectable as visualized by the green fluorescein isothiocyanate fluorescence.

ICAM-1 was strongly and constantly expressed by the chondrocytes in the control and VIB group. ICAM-1 was highly accumulated around the nucleus and spread out through the cell in both the control and vibration images. No difference was detected between both groups (Fig. 2N, Q).

**Table 2.** Primers used for quantitative real-time PCR. All sequences are given in 5'-3'-direction

Gene	Direction	Sequence
18S rRNA	forward	GGAGCCTGGCGCTTAATTT
	reverse	CAACTAAGAAGCCGCATGCA
ACAN	forward	AGTCCAACCTTCAAGGTGAAC
	reverse	ACTCAGCGAGTTGTCATGGT
ANXA1	forward	GCCAAAGACATAACCTCAGACACAT
	reverse	GAATCAGCCAAGTCTTCATTGACA
ANXA2	forward	GGTACAAGAGTTACAGCCCTTATGACA
	reverse	CATGGAGTCATACAGCCGATCA
BAX	forward	GTCAGCTGCCACTCGGAAA
	reverse	AGTAACATGGAGCTCGAGAGGAT
BCL2	forward	TCAGAGACAGCCAGGAAAATCA
	reverse	CCTGTGGATGACTGAGTACCTGAA
BIRC2	forward	GCTTTTGTGTGATGGTGGCT
	reverse	ACTCACACCTTGGAAACCACT
BIRC3	forward	TGCTGTGATGGTGACTCAG
	reverse	ACTCACACCTTGGAAACCACT
BIRC5	forward	GCCAGATGAGACCCCATAG
	reverse	CACCAAGGGTTAATCTTCAAACCTG
COL1A1	forward	ACGAAGACATCCCACCAATCAC
	reverse	CGTTGTGCGAGACGAGAT
COL2A1	forward	GGCAATAGCAGGTTACAGTACA
	reverse	CGATAACAGTCTTGCCCACTT
CYC1	forward	CACTGCGGGAAGGTCTCTAC
	reverse	GGGGTGCATCGTCAAACCTC
FADD	forward	CCTGGGGAAGAAGACCTGTGTG
	reverse	TGATGCTGCGATCTTGGTG
FAS	forward	AGTCTGGTTTATCCCATTTGAC
	reverse	AGGGATTGGAATTGAGGAAGACT
FN1	forward	TGAGGAGCATGTTTTAGGAGAA
	reverse	TCCTCATTACATCGGCGTATAC
NFKB1	forward	CTTAGGAGGAGAGCCAC
	reverse	TGAAACATTTGTTCCAGGCCTTC
PXN	forward	CATGGACGACCTCGACGC
	reverse	CAAGAACACAGCCGTTTGG
RELA	forward	CGTTTTCACACACTGGATTTC
	reverse	ACTGCGGATGGCTTCT
TLN1	forward	GATGGCTATTACTCAGTACAGACAAGTGA
	reverse	CATAGTAGACTCTCATCTCTTCCA
VCL	forward	GTCTCGGCTGCTCGTATCTT
	reverse	GTCCACCAGCCCTGTCATTT

**Table 3.** Primary antibodies for Western blotting. Ms: mouse, Rb: rabbit

Antibodies	Company / No	Species	Detected MW (kDa)	Dilution
Anti-β-actin	Sigma / #A5316	Ms	42	1:5000
Anti-Caspase 3	Cell-Signaling / #9662S	Rb	17, 19, 35	1:800
Anti-NFκB p65	Cell-Signaling / #4764S	Rb	65	1:800

*Expression of selected genes*

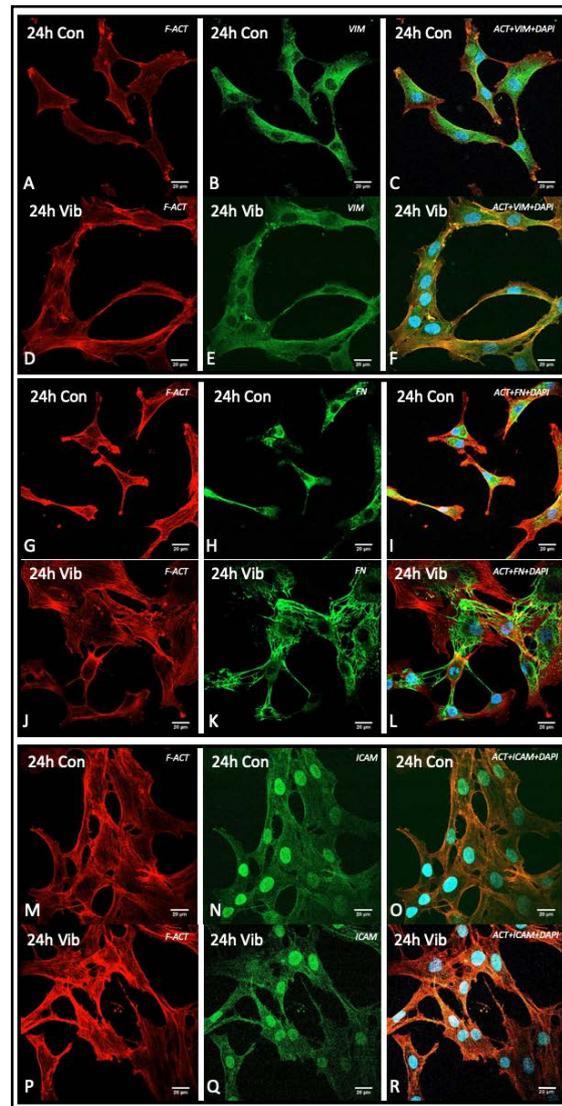
In order to examine the influence of VIB on cultured chondrocytes, the gene expression of selected genes belonging to biological processes such as apoptosis, extracellular matrix, cell adhesion, or focal adhesions were studied. The chosen set of 19 genes comprised *ACAN*, *COL1A1*, *COL2A1*, *FN1* (Fig. 3), *VCL*, *TLN1*, *PXN* (Fig. 4), and *ANXA1*, *ANXA2*, *NFKB1*, *RELA*, *BIRC2*, *BIRC3*, *BIRC5*, *FADD*, *FAS*, *CYC*, *BAX*, *BCL2* (Fig. 5).

Significant changes were found for *VCL*, *PXN*, *ANXA1*, *ANXA2*, *BAX* and *BCL2* mRNAs. All others were not significantly altered (Fig. 3-5). Although the gene expression of *RELA* was not changed after a 24-hour vibration-exposure, the NF- $\kappa$ B p65 protein was down-regulated in VIB samples, which hints to the cell-protective effects of low frequency VIB on human chondrocytes *in vitro* (Fig. 5).

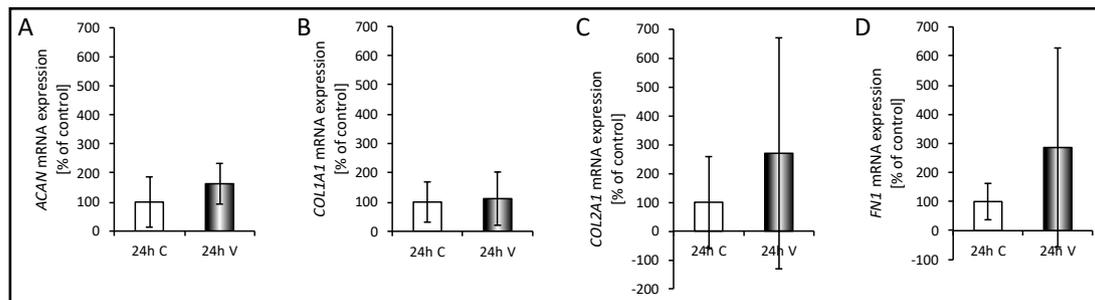
*Semantic analysis of observed gene and protein alterations in combination with earlier data*

As determined by qPCR the majority of the 19 genes remained unchanged after 24 h of treatment (Fig. 3-5). Only 6 genes (*VCL*, *PXN*, *ANXA1*, *ANXA2*, *BAX* and *BCL2*) were significantly changed in the chondrocytes of the six donors after 24 h of incubation on a Vibraplex (Fig. 3-5). In order to estimate the relevance of this data, we compared them with earlier data shown in Tables 4-6 using the Pathway Studio program.

We did not find a significant up-regulation of *ACAN* and *COL2A1* (Fig. 3). According to the pathway analysis shown in Fig. 6, which is based on a number of

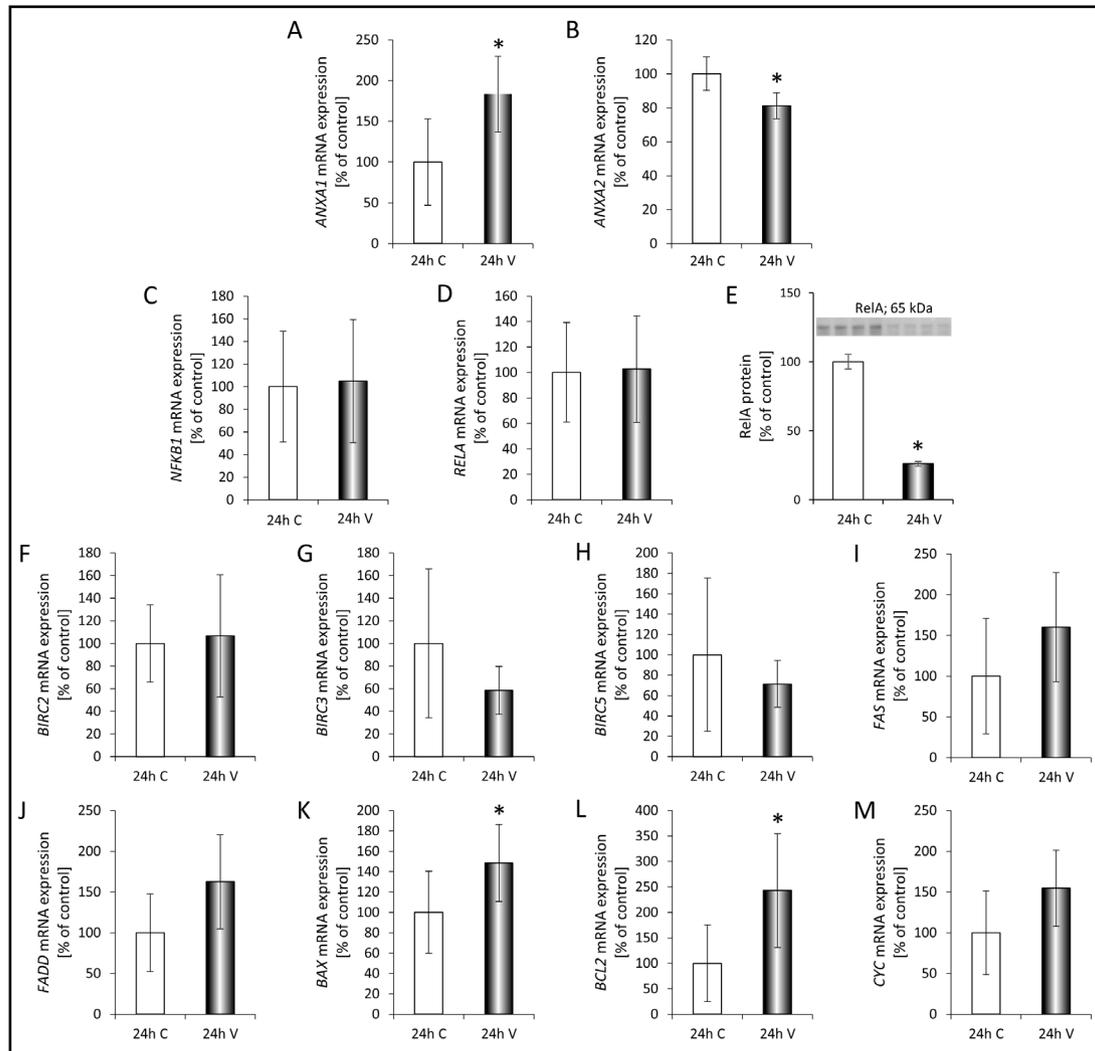
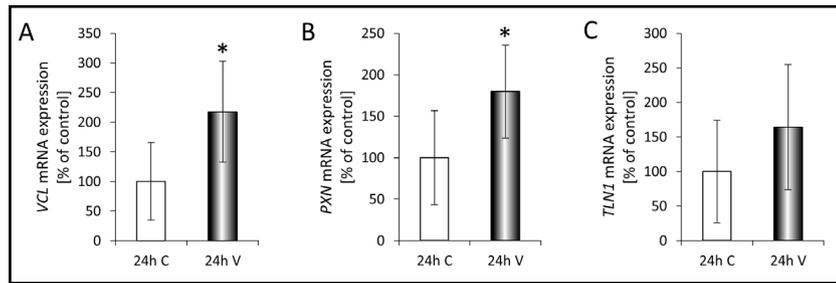


**Fig. 2.** Immunofluorescence staining of vimentin (B, E), fibronectin (H, K) and ICAM-1 (N, Q) (green), F-actin filaments (red; A, D, G, J, M, P) and the nucleus (blue - DAPI) in chondrocytes exposed to static control conditions (A-C, G-I, M-O) or to VIB (D-F, J-L, P-R) for 24 h. The scale bar is 20  $\mu$ m.



**Fig. 3.** Gene expression of ECM genes were determined by qPCR. Data is presented as means  $\pm$  standard deviation; n=6 (qPCR); C = control; V = VIB; \* = p<0.05.

**Fig. 4.** Gene expression of focal adhesion genes were determined by qPCR. Data is presented as means  $\pm$  standard deviation; n=6 (qPCR); C = control; V = VIB; \* = p<0.05.



**Fig. 5.** Gene expression of the intrinsic and extrinsic pathway of apoptosis were determined by qPCR. In addition, the NF- $\kappa$ B-p65 protein content was also analyzed by Western blot. Data is presented as means  $\pm$  standard deviation; n=6 (qPCR) and n=4 (Western blot); C = control; V = VIB; \* = p<0.05.

studies described in the literature, an up-regulation of *COL2A1* may be supported by *ACAN*, as observed in [15] and by *RELA*. *COL2A1* may induce an up-regulation of *CXCL8* and *CASP3*, whereas *IL6* inhibits *COL2A1* expression.

According to [3], *ANXA2* was down-regulated while *SOX9* was up-regulated. *ANXA2* inhibits *CASP3* accumulation (Fig. 7). In addition, *COL2A1* inhibits *ACTB*. Enhanced *SOX5* and *SOX9* transcription factors are regulating the actin cytoskeleton (Fig. 7). In Fig. 3 we demonstrate that *COL2A1* was unchanged by VIB, but *COL1A1* was unchanged and expressed

at a low level. *ACTB* inhibits *SOX9* expression and activates *ACAN*, (Fig. 6) as it is described in [16].

In an earlier study, *SPP1* was also enhanced after 31P at gene and at protein levels [3]. At that time *ACTB*, *CAV1*, *ICAM1*, *SOX9*, *TUBB* and *VIM* were up-regulated together with *SPP1*, while *SOX5* was down-regulated. This may be explained by a mutual up-regulation of *SPP1*, *SOX9*, *VIM*, and *ICAM1* (Fig. 6, green arrows) [17-19]. *SPP1* had been shown to positively influence *IL6* and *CXCL8* genes (Fig. 6) [20]. Wehland *et al.* [21] had demonstrated that the *CXCL8* gene expression was up-regulated in vibrated chondrocytes and chondrocytes exposed to 31P during a PF mission (Table 5). They concluded that vibration was the major driving factor inducing the increase in *CXCL8* mRNA [21].

Above, we showed that the *ANXA2* mRNA was significantly down-regulated in the six donors (Fig. 5). Cleaved caspase-3 (17 kDa) was not detected in chondrocytes exposed to VIB for 24 h (Fig. 1G), as well as factors of the extrinsic pathway of apoptosis like FAS and FADD were not altered by vibration. Furthermore, *BAX* and *BCL2* genes were both elevated and thus inhibiting each other (Fig. 8).

Furthermore, *SPP1* could play an inhibitory role on *CASP3* gene expression (Fig. 6, red line with cross bar) [22]. NF- $\kappa$ B p65 has an influence on aggrecan (Fig. 7). Interestingly, aggrecan and collagen type 2 were found enhanced in chondrocytes after 24 h on the RPM [15], but not significantly changed in VIB samples (Fig. 3), although there was a tendency to increase. In addition, after 16 days and 4 months of exposing chondrocytes to (s-)  $\mu$ g, collagen type II was diminished, and collagen type 1 and type 10 were reduced already after 24 h [15, 23, 24]. In (s-)  $\mu$ g, *SPP1* has shown an upregulating influence on the ECM components aggrecan and collagen type 2, but not on collagen type 1 and type 10 (Table 6) [25]. In patients suffering of OA, osteopontin may exert a protective effect against aggrecan degradation [26]. *SOX5* and *SOX9* play essential roles in producing aggrecan and collagen type 2 during cartilage formation [27-29].

**Table 4.** Comparison of gene and protein expression analyses with literature data from earlier experiments. ▼: significantly down-regulated; ○: no regulation; ▲: significantly up-regulated; n.d.: not determined; n.e.: not expressed; PFC: Parabolic flight campaign; RPM: Random Positioning Machine; VIB: Vibration; 31P: after 31 parabolas

Gene/Protein	Gene Expression				Protein Expression	
	VIB [6]	[3]	[3]	[3]	[15]	
	VIB vs. 1g	PFC vs. 1g	VIB vs. 1g	1.8g vs. 1g	PFC 31P vs. 1g	24 h RPM vs. 1g
ACTB	○	▲	○	○	▲	n.d.
ANXA2	▼	n.d.	n.d.	n.d.	n.d.	n.d.
CASP3	○	n.d.	n.d.	n.d.	n.d.	▼
CAV1	○	▲	○	○	n.d.	n.d.
GAPDH	▲	n.d.	n.d.	n.d.	n.d.	n.d.
ICAM1	○	▲	○	○	▲	n.d.
ITGB1	○	n.d.	n.d.	n.d.	n.d.	n.d.
RELA	○	n.d.	n.d.	n.d.	n.d.	n.d.
SOX5	○	▼	○	○	n.d.	n.d.
SOX9	▲	▲	○	○	▲	n.d.
SPP1	○	▲	○	○	▲	n.d.
TUBB	○	▲	○	○	▲	n.d.
VIM	○	▲	○	○	▲	n.d.

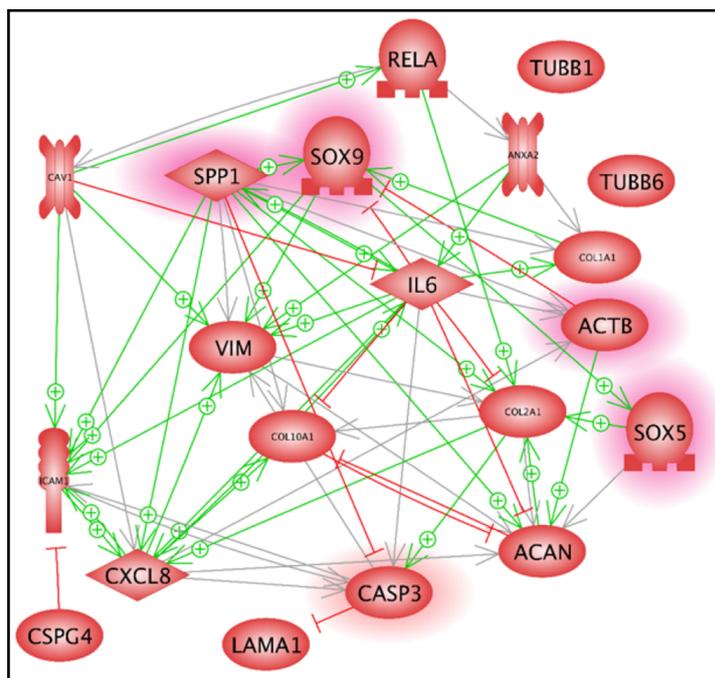
**Table 5.** The gravitational influence on the pro-inflammatory cytokines IL-6 and IL-8 [21]. ▼: significantly down-regulated; ○: no regulation; ▲: significantly up-regulated; n.d.: not determined; PFC: Parabolic flight campaign; VIB: Vibration

Gene/Protein	Gene Expression		Protein Expression	
	VIB 2h vs. 1g	PFC 2h vs. 1g	VIB 2h vs. 1g	1.8g 2h vs. 1g
IL6	▲	○	○	▲
CXCL8	▲	▲	n.d.	n.d.

**Table 6.** Regulation of extracellular matrix components under altered gravity. ▼: significantly down-regulated; ○: no regulation; ▲: significantly up-regulated; n.d.: not determined; PFC: Parabolic flight campaign; RPM: Random Positioning Machine; 31P: after 31 parabolas

Protein/ Gene	Protein Expression				Gene Expression
	[15]	[4]	[23]	[24]	[21]
	24h RPM vs. 1g	24h RPM vs. 1g	16d RPM/r- $\mu$ g vs. 1g	4mo r- $\mu$ g vs. 1g	PFC 31P vs. 1g
Aggrecan	▲	n.d.	▼	n.d.	n.d.
Chondroitin Sulfate	▲	n.d.	n.d.	n.d.	n.d.
Coll. Type I	▼	n.d.	▼	n.d.	n.d.
Coll. Type II	▲	▲	▼	▼	▼
Coll. Type X	▼	n.d.	n.d.	n.d.	n.d.
Laminin	○	n.d.	n.d.	n.d.	n.d.

**Fig. 6.** Mutual interaction of genes and their products on a genetic level. The genes were selected from the studies described in Tables 4-6. The graph was generated with Pathway Studio v.11 software (Elsevier Research Solutions, Amsterdam, The Netherlands).



Therefore, it may be suggested that microgravity induces an up-regulation of *SPP1* genes in chondrocytes and cartilage. The STRING analysis given in Fig. 8 demonstrated a central role for *FN1*, *VCL*, *NFKB1*, *RELA*, *FADD* and *CYC1*.

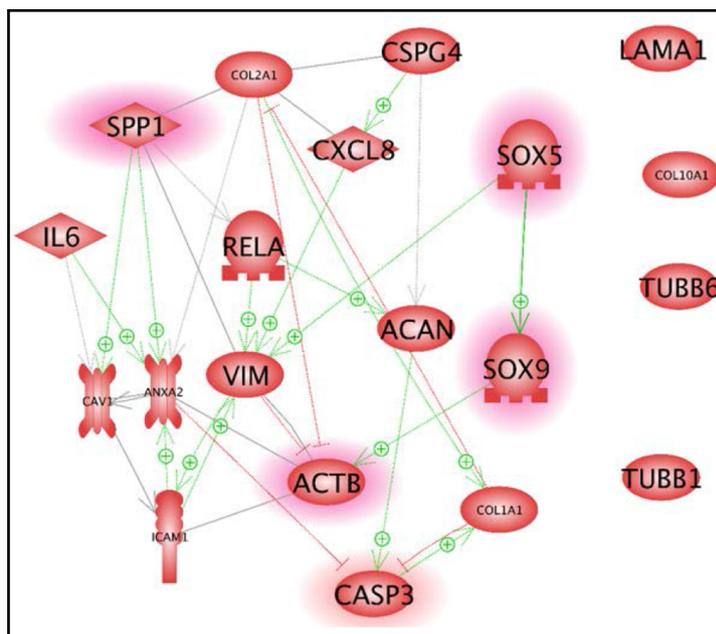
## Discussion

In transportation systems like planes or spacecrafts one source of travelers' or crewmembers' discomfort are VIBs. They are experienced by the passengers due to the motion of the aircraft or the spacecraft in its operational mode. These VIBs can occur during normal passenger flights, but also during PF and space missions. Studies applying WBV have demonstrated an influence on bone and cartilage and propose WBV for the prevention of subchondral bone loss of knee OA [8]. Little is known about the direct effects of low frequency VIB on human chondrocytes, occurring during a long-term passenger flight. Therefore, we focused on the impact of low frequency VIB, measurable during a normal flight. It is known that short-term  $r\text{-}\mu g$  occurring during PF maneuvers ( $r\text{-}\mu g$ ) for about 2 hours induced only moderate changes in human chondrocytes and a two-hour-exposure to VIB exerted no damage of the cells [5].

Cartilage tissue has a low regeneration potential. The chondrocytes show a well-differentiated phenotype with unique physiological functions. Chondrocytes have shown to react to external stress and forces *in vitro* [3-6]. For example, culturing chondrocytes in simulated microgravity using  $\mu g$ -simulation devices like the NASA-developed rotating wall vessel (RWV) or the RPM induces three-dimensional growth of the chondrocytes and favor cartilage formation [30, 31].

However, long-term space missions revealed bone loss and cartilage breakdown in crewmembers, resulting in similar symptoms compared to OA [32, 33]. Therefore, it is of high interest to design and perform new studies focusing on chondrocytes *in vitro* or cartilage *in vivo*. Recently, a paper demonstrated the effects of low-magnitude WBV on cartilage degradation, bone/cartilage turnover, and OA joint function in a rabbit knee OA model [33, 34]. Lower frequency (20 Hz) WBV revealed beneficial effects for the bone micro-structure [34].

**Fig. 7.** Mutual interaction of proteins, which were selected from the studies described in the Tables 4-6. The graph was generated with Pathway Studio v.11 software (Elsevier Research Solutions, Amsterdam, The Netherlands).



We used human chondrocytes deriving from six different donors. The cells were exposed to the Vibraplex device *in vitro*. We did not detect any VIB-induced morphological changes and no cell death, which confirmed earlier data [6]. Recently, we had performed a whole genome microarray analysis on vibrated chondrocytes. Overall, we detected that VIB had only very little influence on the differential gene expression in chondrocytes [6].

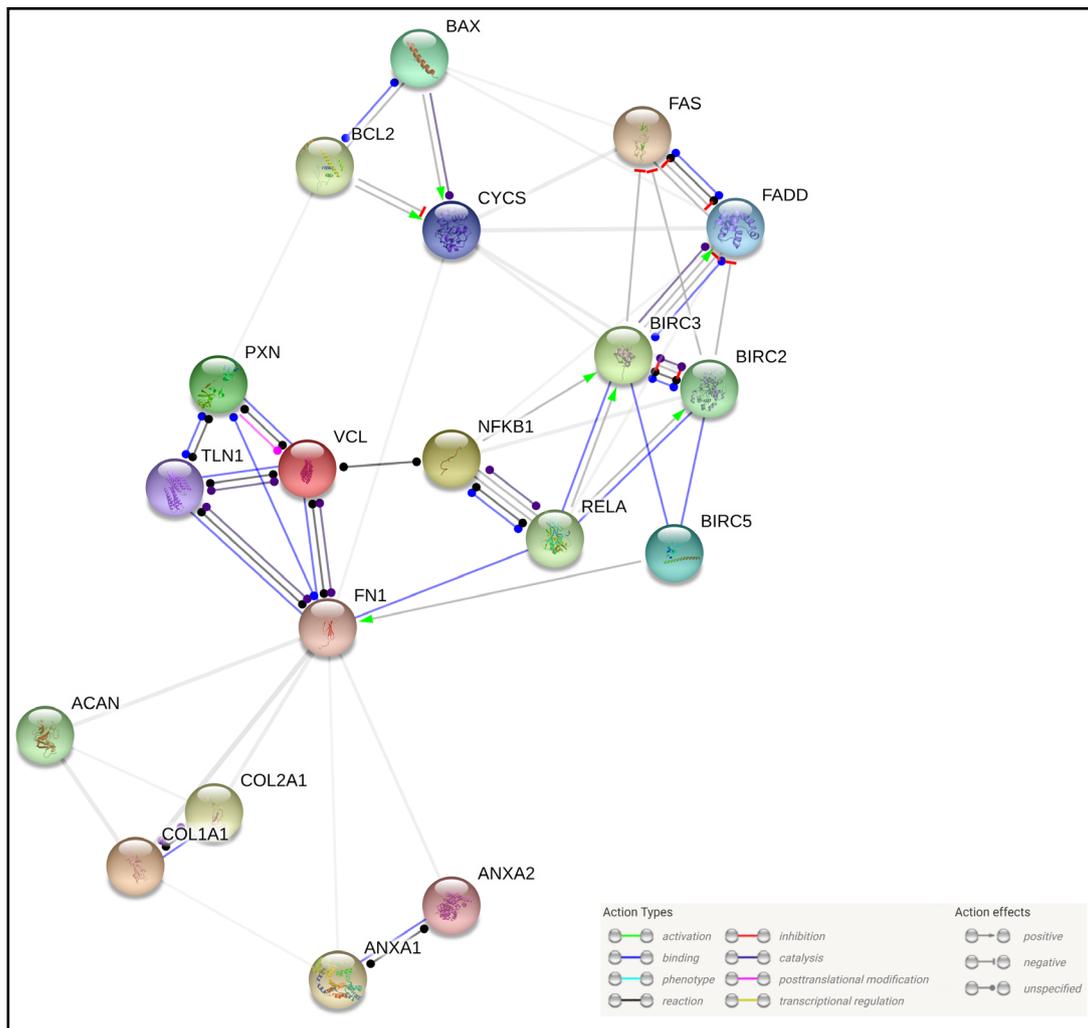
In this study, we focused on selected genes belonging to the extracellular matrix which are characteristic for chondrocytes (*ACAN*, *COL2A1*), on focal adhesion markers and on factors of the intrinsic and extrinsic pathway of apoptosis. The objective was to further verify our hypothesis that VIB might have beneficial effects on chondrocytes and eventually on cartilage.

Long-term VIB of human chondrocytes *in vitro* promotes the interaction of cells leading to changes in extracellular matrix proteins or cell adhesion molecules such as fibronectin. Fibronectin was disorganized and accumulated in the extracellular space in vibrated chondrocytes compared to control cells (Fig. 2). Fibronectin is a key factor of the osteoarthritic process and it plays a central role in this study as demonstrated by the interaction network shown in Fig. 8. In particular, in the early stages of OA fibronectin has shown a rapid accumulation of the ECM, which might suggest an attempt of the tissue to repair itself [34]. A large elevation of *FN1* and *SPP1* mRNAs was measured in human OA-affected cartilage samples as compared with normal cartilage [35, 36]. Both cell adhesion molecules are involved in the inflammatory process of OA.

*SPP1* was not significantly altered by vibration which hints to a more beneficial effect [6, 35], as an elevated osteopontin content could be involved in cartilage damage in astronauts, where osteopontin is linked to severity and progression of OA. Studies have shown that OA patients present higher concentrations of osteopontin as compared to healthy patients [3, 4, 8, 26, 37, 38].

We investigated whether focal adhesion genes are influenced by vibration in chondrocytes (Fig. 3-5). Focal adhesion proteins link the ECM and the actin cytoskeleton. They propagate signals arising from the activation of integrins following their engagement with ECM proteins like fibronectin. Fibronectin binding by cells influences vinculin (Fig. 8). Vibration significantly elevated the *VCL* mRNA expression. A fibronectin-vinculin transmembrane interaction is involved in differentiation processes [39].

Midpalatal suture cartilage cells exposed to expansive stimulation form actin stress fibers, and a reorganization of focal adhesion contacts (vinculin) occurred [40]. In parallel,



**Fig. 8.** STRING analysis with the major protein-protein interactions for chondrocytes experiments under mechanical vibration. The visualization was made using the online software STRING [https://string-db.org/].

fibronectin was elevated [39], which is comparable to our *in vitro* findings. Vibration also elevated the *PXN* mRNA in human chondrocytes (Fig. 4). The association of focal adhesion kinase with fibronectin and paxillin is required for precartilaginous condensation of chick mesenchymal cells [41]. Paxillin serves as a critical transducer of signals from fibronectin [42]. Vinculin has shown to regulate the survival of mouse embryonal cancer cells through regulating paxillin-FAK interactions to alter ERK1/2 activation [43]. In addition, paxillin is acting anti-apoptotic and is involved in the survival of mouse cells [44].

We have focused on apoptosis applying TUNEL staining. In all cultures of the six donors no morphological signs of programmed cell death were detectable. The *CASP3* and *RELA* gene expression was not significantly changed, but a significant reduction of the *ANXA2* gene expression was found in vibrated chondrocytes after 24 h. Cleaved caspase-3 protein (17 kDa) was not detectable. As the chondrocytes are the only cell type found in cartilage, inhibition of programmed cell death may be crucial for the maintenance of healthy cartilage [45]. The protein of *ANXA2* is annexin-A2, which is belonging to the annexin family. It plays a role in the development of OA. Annexin-A2 is involved in regulating cellular growth, cell motility and in signal transduction pathways (extracellular matrix). Human osteoarthritic chondrocytes undergo terminal differentiation, and are releasing, among other factors,

annexin A II-containing matrix vesicles which are able to initiate mineral formation, and ultimately die by apoptosis. Thus, these cells are involved in the destruction of articular cartilage in OA [46]. Moreover, we detected an up-regulation of *ANXA1* mRNA after vibration. *ANXA1* inhibits the activation of NF- $\kappa$ B (nuclear factor  $\kappa$ -light-chain-enhancer of activated B cells) by binding to the p65 subunit [47], it is anti-proliferative, anti-apoptotic and associated with survival in gastric cancer [48].

In this study the gene expression of *RELA* was not changed after a 24-hour vibration-exposure, but the NF- $\kappa$ B p65 protein was down-regulated in VIB samples, which hints to the cell-protective effects of low frequency VIB on human chondrocytes *in vitro*. In addition, we studied key factors of the intrinsic and extrinsic pathway of programmed cell death. *BAX* (Bcl-2 associated X-protein) and *BCL2* mRNAs were significantly up-regulated, whereas *CYC1* was not changed. In addition, *FAS* and *FADD* were not changed by vibration. In a next step, we investigated the *BIRC2*, *BIRC3*, *BIRC5*, which are members of the protein (IAP) family. IAPs are regulating the caspase activity, as well as proliferation and survival. None of these genes were changed in expression, neither *BIRC2*, which is the gene of the Baculoviral IAP repeat-containing protein 2, nor *BIRC3*, which inhibit apoptosis by binding to the tumor necrosis factor receptor-associated factors TRAF1 and TRAF2. *BIRC2* inhibits the activation of caspases-3, -7 and -9 [49]. *BIRC5* gene is coding for the surviving factor was also not changed by vibration. Hence, we did not find signs that vibration for 24h modulates chondrocyte apoptosis.

These experiments demonstrated that long-term VIB showed beneficial effects. The up-regulation of *VCL*, *PXN*, *ANXA1* and the down-regulation of *ANXA2* seem to protect the chondrocytes and inhibits dedifferentiation.

## Conclusion

In conclusion, a 24-hour exposure to vibration as generated by the Vibraplex showed beneficial effects on human chondrocytes which did not reveal any noteworthy morphological alterations or damage. Pathway analyses hint towards a beneficial influence of this kind of vibration on human cartilage cells. The future of space exploration will involve long-duration spaceflights and stays on the International Space Station or flights to the Mars and beyond, and crewmembers will be subjected to a number of stressors like microgravity, vibration or cosmic radiation. Therefore, long-term vibration and microgravity studies on different cell types like endothelial cells [50] or lymphocytes [51] should be performed in the future.

## Abbreviations

$\mu$ g (Microgravity); 3D (Three dimensional); ACAN (Aggrecan); ACTB (Beta-actin); *ANXA1* (Annexin a1); *ANXA2* (Annexin a2); *BAX* (Apoptosis regulator BAX); *BCL2* (Apoptosis regulator BCL2); *BIRC2* (Baculoviral IAP repeat-containing protein 2); *BIRC3* (Baculoviral IAP repeat-containing protein 3); *BIRC5* (Baculoviral IAP repeat-containing protein 5); BSA (Bovine serum albumin); *CASP3* (Caspase-3); *COL1A1* (Collagen alpha-1(I) chain); *COL2A1* (Collagen alpha-1(II) chain); CON (Control); *CXCL8* (Interleukin-8); *CYC1* (Cytochrome c1); DAPI (4',6-diamidino-2-phenylindole); DLR (Deutsches Zentrum für Luft- und Raumfahrt); DPBS (Dulbeccos phosphate buffered saline); ECM (Extracellular matrix); ERK1/2 (Mitogen-activated protein kinase 3/6); *FADD* (FAS-associated death domain protein); *FAS* (Tumor necrosis factor receptor superfamily member 6); FITC (Fluorescein isothiocyanate); FN1 (Fibronectin 1); IAP (Inhibitor of apoptosis protein); ICAM-1 (Intercellular adhesion molecule 1); IL6 (Interleukin-6); NF- $\kappa$ B p65 (Nuclear factor NF-kappa-B p65 subunit); NFKB1 (Nuclear factor NF-kappa-B p105 subunit); OA (Osteoarthritis); PCR (Polymerase chain reaction); PF (Parabolic flight); PSMD4 (26S proteasome non-ATPase regulatory subunit 4); PVDF (Polyvinylidene fluoride); *PXN* (Paxillin); r- $\mu$ g (real microgravity); *RELA*

(Nuclear factor NF-kappa-B p65 subunit); RPM (Random positioning machine); RWV (Rotating wall vessel); s- $\mu$ g (simulated microgravity); SOX5 (Transcription factor SOX-5); SOX9 (Transcription factor SOX-9); SPP1 (Osteopontin); TBX15 (T-box transcription factor TBX15); TRAF 1/2 (TNF receptor-associated factor 1/2); TUBB (Beta-tubulin); TUNEL (TdT-mediated dUTP-biotin nick end labeling); VCL (Vinculin); VIB (Vibration); VIM (Vimentin); WBV (Whole body vibration).

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## Disclosure Statement

The authors declare no competing financial interests.

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