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Tuning Mesenchymal Stem Cell Response onto Titanium-Niobium-Hafnium Alloy by Recombinant Fibronectin Fragments

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ABSTRACT

Since metallic biomaterials used for bone replacement possess low bioactivity, the use of cell adhesive moieties is a common strategy to improve cellular response onto these surfaces. In recent years the use of recombinant proteins has emerged as an alternative to native proteins and short peptides owing to the fact that they retain the biological potency of native proteins, while improving their stability. In the present study, we investigated the biological effect of two different recombinant fragments of fibronectin, spanning the 8th-10th and 12th-14th type III repeats, covalently attached to a new TiNbHf alloy using APTES silanization. The fragments were studied separately and mixed at different concentrations and compared to a linear RGD, a cyclic RGD and the full-length fibronectin protein. Cell culture studies using rat mesenchymal stem cells demonstrated that low to medium concentrations (30% and 50%) of type III 8th-10th fragment mixed with type III 12th-14th fragment stimulated cell adhesion and proliferation compared to RGD peptides and the fragments separately. On the other hand, type III 12th-14th fragment alone or mixed at low volume percentages $\leq 50\%$ with type III 8th-10th fragment increased alkaline phosphatase levels compared to the other molecules. These results are significant for the understanding of the role of fibronectin recombinant fragments in cell responses and thus to design bioactive coatings for biomedical applications.

KEYWORDS: Fibronectin; recombinant protein; cell adhesive peptides; mesenchymal stem cells; titanium alloy

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1. INTRODUCTION

Titanium (Ti) and Ti-based alloys have been traditionally used for bone implantation purposes.[1] Despite the good biocompatible properties showed by these materials, there is a mismatch with bone elastic modulus, which is translated in a stress shielding effect.[2] In a bone replacement scenario, the prosthesis shares the load and carrying capacity of bone. The loads applied to bone are reduced leading to a new loads distribution and eventually to bone resorption, which is a major cause of implant failure. Ductile metals such as Zr, Ta, Nb or Hf are used to stabilize the beta phase in Ti-alloys,[3–5] which is the phase with lowest elastic modulus, and also because they possess osseoinductive properties.[6] A particularly interesting Ti-based alloy is the TiNbHf ternary system. This system has shown interesting properties such as low elastic modulus, good biocompatibility and no cytotoxic effects.[7,8] In this regard, we have recently developed the Ti25Nb21Hf alloy, which exhibited lower elastic modulus and higher wear resistance compared to Ti.[9]

Despite the interesting biocompatibility and mechanical properties exhibited by TiNbHf alloys, these metallic biomaterials are not bioactive and thus do not promote osseointegration at an optimal rate, compromising in some circumstances the long-lasting stability of the implants. For this reason, in recent years several efforts have concentrated on the biofunctionalization of the biomaterials surface to improve their bioactivity and hence their osseointegration.[10] In detail, the main goal of biofunctionalization is to generate an extracellular matrix (ECM)-like environment that positively influences the behaviour of osteoblast and bone precursors (i.e. mesenchymal stem cells, MSCs) and effectively guides bone growth.[11,12] The ideal strategy is the use of native bone ECM proteins, such as collagen or fibronectin (FN), for coating the surfaces.[13,14]

However, although these proteins retain their full biological functionality and specificity, they display low stability after functionalization along with many other drawbacks including

high production costs and immunogenicity, which have reduced their biomedical potential.[11]

The use of ECM-derived synthetic peptides containing the functional sites of ECM proteins is

an interesting approach to overcome these problems.[10,11] The advantages of these motives in

regards to ECM proteins are their higher stability, lower production costs and their capability

to be synthetically tailored in composition for specific biological applications.

In particular, integrin-binding sequences are of great interest since they regulate not only cell

adhesion but also cell survival, proliferation and differentiation. This is illustrated by the cell

adhesive arginine-glycine-aspartic acid (RGD) motif, which is present in many ECM proteins

including FN, vitronectin, bone sialoprotein and osteopontin, and has been widely used to

mimic cell/matrix interactions and improve the osteointegration of implant materials.[10–12,15]

Nonetheless, short synthetic peptides are commonly less functional and specific than full-

length proteins due to the absence of complementary domains required for biological activity

or inadequate conformation, both of which are crucial to trigger an appropriate cell

response.[16] To solve these problems, different approaches aiming at improving the activity

and selectivity profile of integrin ligands have been followed: cyclic peptides,[17–19]

peptidomimetics,[20–23] mixtures of peptides[24–26] or peptide-based platforms.[27]

The use of recombinant protein fragments for functionalizing surfaces has emerged as a

potential strategy that combines the benefits of the aforementioned methods.[28–31] Using

recombinant DNA technology the signalling domain of an ECM protein can be engineered to

retain crucial biological features such as the conformation and spacing of bioactive

sequences, while reducing the antigenicity and improving the stability of native proteins.

Noteworthy, fusion of more than one domain from the same protein or domains from

different proteins can be assembled modulating different cell responses with only one

molecule.[32–34]

FN is a well-known ECM protein that mediates many cellular processes like cell adhesion, migration, growth and differentiation.[35] FN contains a region involved in cell adhesion located in the type III_{10} domain, which includes the above-mentioned RGD sequence. In addition, the type III_9 domain contains the PHSRN (Pro–His–Ser–Arg–Asp) sequence, which strongly modulates and increases the binding affinity of RGD to $\alpha5\beta1$ integrin.[35,36]

Previous studies have demonstrated that the cell attachment site (CAS) recombinant fragment of FN, consisting of type III_{8-10} repeats or III_{7-10} enhances cell adhesion, proliferation and differentiation.[28,37] Studies by Martino et al.[31] demonstrated the good adhesion and osseoinduction potential of MSCs in contact with the $FNIII_{9-10}$ fragment. Petrie et al.[30] showed the capacity of the FN III_{7-10} fragment to increase osteoblastic differentiation and mineralization *in vitro* and enhanced implant osseointegration in a rat cortical bone model compared to RGD and fibronectin. In vivo studies by Agarwal et al. demonstrated that the FN III_{7-10} fragment promotes $\alpha_5\beta_1$ integrin-dependent adhesion and osteogenic differentiation of hMSCs.[38]

Besides the CAS, there is another region in FN, the C-terminal type III_{12-14} heparin-binding II (HBII) domain also crucially involved in cytoskeletal organization and focal adhesion formation. This HBII domain binds to proteoglycans, including heparan sulfate proteoglycans (HSPGs) such as syndecans.[39] For instance, syndecan-4 has been described to play an important role in the regulation of the activation of several integrins through the HBII domain.[40] Moreover, it has been shown that HB domains in FN enhance cell adhesion and proliferation of osteoblasts together with the CAS domain.[39] Another important feature of the HBII domain was described by Martino et al.[41] In his studies, its capability to sequester growth factors from different families was demonstrated.

The present work aims to covalently functionalize a Ti25Nb21Hf alloy with FNIII $_{8-10}$ and FNIII $_{12-14}$ recombinant fragments alone or combined at different concentrations to elucidate

the importance of each fragment on the biological response for implant applications. These sequences have been compared to FN and two short synthetic RGD peptides (linear and cyclic, linRGD and cRGD, respectively). Cellular behaviour has been analysed in terms of cell adhesion, proliferation and differentiation using rat mesenchymal stem cells (rMSCs).

2. MATERIALS AND METHODS

2.1. TiNbHf alloy fabrication and preparation

The alloy was fabricated as described in [9]. Briefly, commercially pure (CP)-Ti, Nb (99.8% purity) and Hf (99.7% purity) were arc melted and vacuum homogenized. Bars of diameter 10 mm were obtained by extrusion. TiNbHf alloy bars were cut in discs of 2 mm thickness and grinded with SiC papers of grit 320, 800, 1200 and 2500 (Struers, Spain). After grinding, the discs were mirror polished with colloidal silica (size particle 0.05 μm, ATM GmbH, Germany). Samples were cleaned in an ultrasound bath with cyclohexane, isopropanol, ethanol, deionized water and acetone (all chemicals were of the highest purity available from Sigma-Aldrich, USA).

2.2. Recombinant FN fragments

Recombinant DNA techniques were used to synthetize two different fragments of human FN: i) the Cell Attachment Site (CAS) sequence, spanning the 8th–10th type III repeats, and ii) the Heparin Binding II (HBII) sequence spanning the 12th–14th type III repeats. DNA extracted from SaOS-2 cells was specifically amplified by PCR using 5'-GTCGACAAGTTCCTCCTCCCACTGACCT-3' and 5'-GCGGCCGCTTAATGGAAATTGGCTTGCTG-3' (for CAS domain) or 5'-GTCGACAAGCTATTCCTGCACCAACTGA-3' and 5'-GCGGCCGCTGTCTTTTTCCTTCCAATCAGG-3' (for HBII domain). Each DNA fragment was digested with SalI and NotI (New England Biolabs, USA) and ligated into a pGEX-6P-1

plasmid (GE Healthcare, UK). The resulting constructs, containing CAS or HBII with a

glutathione S-transferase (GST) tagging sequence at the amine terminus, were separately

amplified in DH5α cells (Invitrogen, USA), purified and sequenced. Then, BL21 *E.coli*

strains (Invitrogen) were separately transformed with the plasmids and streaked onto LB agar

plates containing 100 μg/ml of ampicillin and incubated at 37 ºC overnight. Colonies were

isolated and dynamically cultured in LB broth (Sigma-Aldrich) with 100 μg/ml of ampicillin

at 37 ºC until they reached an OD_{600}=0.6. Expression of the fragments was induced by

addition of 1 mM IPTG. After 4 h, cells were harvested by centrifugation and lysed by

sonication. Then, 20% Triton X-100 was added and the suspension was incubated at 4 ºC for

30 min under mild agitation. The suspension was then centrifuged and the supernatant was

purified at 4 °C using GSTrap affinity columns (GE Healthcare) in an ÄKTA purifier (GE

Healthcare). GST tag was on-column removed by cleavage with HRV3C Protease (Sigma-

Aldrich) in Cleavage Buffer (50 mM Tris-HCl, 150 mM NaCl, 1 mM EDTA, 1 mM

dithiothreitol, pH 7.0). Purity was verified to be >95% by SDS-PAGE. Protein concentrations

were determined by BCA assay (Pierce, Thermo Fisher Scientific, USA).

2.3. Synthesis of linear RGD peptide (linRGD)

The linear peptide MPA-Ahx-Ahx-Ahx-Gly-Arg-Gly-Asp-Ser-OH (**linRGD**) (Ahx:

aminohexanoic acid; MPA: 3-mercaptopropionic acid) was manually synthesized by solid-

phase peptide synthesis (SPPS) following the Fmoc/tBu strategy[42] and using 2-chlorotrityl

chloride resin (CTC, 200 mg, 1.6 mmol/g) (Iris Biotech GmbH, Germany) as solid support.

Fmoc-L-amino acids were obtained from Iris Biotech GmbH and coupling reagents from

Sigma-Aldrich and Luxembourg Industries Ltd. (Israel). All other chemicals and solvents

were acquired from Sigma-Aldrich, Alfa Aesar (Germany) and SDS (France). The first amino

acid was incorporated by sequential addition of Fmoc-Ser(tBu)-OH (1 equiv) and N,N-

diisopropylethylamine (DIEA) (10 equiv). After 1 h under stirring the resin was capped with

MeOH (0.8 mL/g of resin). The Fmoc group was removed by treatment with piperidine/N,N-dimethylformamide (DMF) (1:4, v/v) and the following coupling reactions were carried out with Fmoc-L-amino acids (4 equiv), Oxyma Pure (4 equiv) and N,N'-diisopropylcarbodiimide (DIC) (4 equiv) in DMF for 45 min. The efficiency of each reaction was monitored using the Kaiser test and by analytical HPLC analysis (Waters Alliance 2695 chromatography system) (Waters, USA). After synthesis, the peptide was cleaved from the solid support with concomitant deprotection of the side chain protecting groups. To this end, the resin was washed with dichloromethane (DCM), dried, and treated with trifluoroacetic acid (TFA)/H_2O/triisopropylsilane (TIS) (95:2.5:2.5) for 1.5 h. The peptide was then precipitated with cold diethyl ether, centrifuged and washed twice with diethyl ether. This crude peptide was dissolved in H_2O/MeCN (1:1, v/v) and lyophilized. Purification of the peptide was achieved by semi-preparative HPLC (Waters Delta 600 instrument) using linear gradients from 0 to 10 % MeCN over 2 min and 10 to 30 % MeCN over 10 min. The purified peptide was characterized by analytical HPLC (10 to 40 % MeCN over 8 min, t_R = 4.227 min) and MALDI-TOF (Applied Biosystems, USA) (m/z calcd. for $C_{38}H_{67}N_{11}O_{13}S$: 917.46, found: 918.30 $[M+H]^+$; 940.28 $[M+Na]^+$); 956.25 $[M+K]^+$).

2.4. Synthesis of cyclic RGD peptide (cRGD)

The cyclic peptide MPA-Ahx-Ahx-Ahx-cyclo(-RGDfK-) (**cRGD**) was synthesized as previously described[19,43,44]. In brief, the thiol-protected anchor MPA(Trt)-Ahx-Ahx-Ahx-OH (1 equiv), 1-hydroxy-7-azabenzotriazole (HOAt) (1.2 equiv) and N-[(dimethylamino)-1H-1,2,3-triazolo[4,5-b]pyridino-1-ylmethylene]-N-methylmethanaminium hexafluorophosphate (HATU) (1.2 equiv) were dissolved in anhydrous DMF (0.022 mM) and pre-activated for 3 h at room temperature in the presence of DIEA (5 equiv). Next, cyclo(-R(Pbf)GD(OtBu)fK-) (1 equiv) dissolved in anhydrous DMF was added to the reaction mixture and allowed to react under stirring overnight at room temperature. After reaction, DMF was evaporated to

dryness, a saturated solution of $NaHCO_3$ was added and the crude product was extracted with EtOAc thrice. The organic layer was washed with brine, dried with Na_2SO_4 and evaporated. The peptide was next treated with TFA/H_2O/TIS (95:2.5:2.5) for 1.5 h. The peptide was finally precipitated with cold anhydrous diethyl ether, washed twice with this solvent and lyophilized. Semi-preparative HPLC purification (linear gradients from 0 to 10 % MeCN over 2 min and 10 to 40 % MeCN over 15 min) yielded the peptide as a white powder. The purified peptide was characterized by analytical HPLC (10 to 40 % MeCN over 8 min, t_R = 6.578 min) and MALDI-TOF (m/z calcd. for $C_{48}H_{78}N_{12}O_{11}S$: 1030.56, found: 1031.39 $[M+H]^+$; 1053.37 $[M+Na]^+$); 1069.35 $[M+K]^+$) .

2.5. Sample biofunctionalization

2.5.1. Silanization

Samples were silanized in order to covalently attach the molecules to the substrate. TiNbHf samples and Quartz Crystal Microbalance with Dissipation Monitoring (QCM-D) Ti sensors (QSX310, Q-Sense, Sweden) were first cleaned in an ultrasound bath with cyclohexane, isopropanol, ethanol, deionized water and acetone, and then cleaned and activated with oxygen plasma for 5 min at a 12 MHz frequency in an Expanded Plasma Cleaner PDC-002 (Harrick Scientific Corporation, USA). Then, samples were immersed in a 0.08 M solution of (3-aminopropyl)triethoxysilane (APTES, Sigma-Aldrich) in toluene (Sigma-Aldrich) at 70 ºC for 1 h under agitation in an inert atmosphere. After silanization, samples were ultrasonicated in toluene for 5 min and cleaned with toluene (3x), acetone (1x), isopropanol (3x), distilled water (3x), ethanol (3x) and acetone (3x). Subsequently aminosilanized samples were immersed in a 7.5 mM solution of N-succinimidyl-3-maleimidepropionate (SMP) in DMF for 1 h under agitation at room temperature. The cross-linked samples were rinsed in DMF (3x), acetone (1x), distilled water (10x), ethanol (3x) and acetone (3x) and dried with nitrogen.

2.5.2. Molecules immobilization

linRGD and cRGD were used at a concentration of 100 μM and CAS and HBII were used at a concentration of 100 μg/ml (concentrations optimized in previous studies in our group). FN (Sigma-Aldrich) was used at a 50 μg/ml concentration. Solvent for all molecules was phosphate buffered saline (PBS, Invitrogen).

Ti25Nb21Hf silanized samples were incubated overnight with a 100 μl drop of linRGD, cRGD, FN, CAS, HBII, respectively, or combinations of different proportions of CAS and HBII (30:70, 50:50 or 70:30 respectively, v/v). After incubation, samples were rinsed in PBS (x3) and blocked with 1% bovine serum albumin (BSA, Sigma-Aldrich) in PBS for 30 min. Samples were sterilized by immersion in ethanol for 30 min followed by 3 rinses in PBS.

2.6. Quartz crystal microbalance with monitoring dissipation (QCM-D)

The QCM-D was used to study the biomolecules adsorption onto silanized Ti sensors. The QCM-D (D300, Q-Sense, Sweden) measurements were performed at 37^0C by monitoring changes in frequency, Δf (Hz), and dissipation, ΔD ($\times 10^{-6}$), in real-time using Qsoft software (Q-Sense). All raw data was analyzed using QTools software (Q-Sense). Frequency and dissipation curves were fitted to a Voigt viscoelastic model to yield relevant mass, thickness, and kinetic information. The description of the Voigt model and details on its implementation using a QCM-D are reported elsewhere.[45,46]

Monitoring of the adsorption was conducted first by completely stabilizing the baseline with PBS for 30-60 min. Then, the biomolecule of study was introduced at a concentration specified in section 2.5.2, and maintained in the sensor chamber for 60 min. Finally, the biomolecules weakly bound to the surface were rinsed with PBS for 10 min.

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2.7. X-ray photoelectron spectroscopy (XPS)

XPS measurements were used to analyze the chemical composition of the samples after biomolecule immobilization. To this end, samples were silanized and coated as explained in section 2.5, and analyzed using an XPS system (SPECS Surface Nano Analysis GmbH, Berlin, Germany) equipped with a Mg anode XR50 source operating at 150 W and a Phoibos 150 MCD-9 detector. Detector pass energy was fixed at 25 eV with 0.1 eV steps to record high resolution spectra at a pressure below 7.5×10^{-9} mbar. Data was analyzed using CasaXPS software (Version 2.3.16, Casa Software Ltd., Teignmouth, UK). Binding energies were calibrated with the C1s signal located at 284.8 eV.

2.8. Cell culture

rMSCs were extracted from femurs of young Lewis rats and expanded in Advanced DMEM supplemented with 10% fetal bovine serum (FBS), 20 mM HEPES buffer solution, penicillin/streptomycin antibiotics (50 U/ml and 50 μg/ml, respectively), 2 mM L-glutamine (all from Invitrogen) at 37 ºC with 5% CO_2 and 95% relative humidity. Cells from passage 5 were used in all the experiments.

2.9. Cell adhesion

Cells (10.000 cells/sample) were seeded in each sample with serum-free medium and allowed to adhere for 4 h. Then, cells were rinsed with PBS (x3) and lysed with 300 μl of Mammalian Protein Extraction Reagent (M-PER, Thermo Fisher, USA). The number of cells adhered was determined using the Cytotoxicity Detection Kit [PLUS] (LDH) (Roche, USA) following the manufacturer's instructions. The lactate dehydrogenase (LDH) activity was measured spectrophotometrically at 492 nm in a PowerWave HT microplate reader (Bio-Tek, USA). A calibration curve with decreasing number of cells was performed to express the results as cell number.

2.10. Cell spreading

Cells (25.000 cells/sample) were seeded in each sample with serum-free medium and fixed 4 h after seeding with 4% paraformaldehyde (Sigma-Aldrich) for 30 min. Fixed cells were permeabilized with 0.05% Triton X-100 in PBS at room temperature for 20 min and washed with 20 mM glycine in PBS (x3). Then samples were blocked with 1% BSA in PBS and incubated for 30 min. Next, cells were incubated with mouse anti-vinculin (1:100; Invitrogen) for 1h and rinsed with 20 mM glycine in PBS. Afterwards, samples were incubated with Alexa Fluor ® 488 goat anti-mouse antibody (1:1000; Invitrogen) and TRITC-phalloidin (1:300; Invitrogen) in the dark for 1 h. After incubation, samples were washed with 20 mM glycine in PBS (x3) and nuclei were counterstained with DAPI (1:1000; Invitrogen). Finally, samples were mounted in Mowiol® 4-88 (Sigma-Aldrich) before visualizing in an E600 fluorescence microscope (Nikon Corp., Japan). Five images at 10X magnification from different areas of each sample were acquired and cell spreading was determined measuring the area of the cells using ImageJ software (National Institute of Health, USA).

2.11. Cell proliferation

Cells were seeded in serum-free medium at a density of 10.000 cells/sample. After 4 h, medium was aspirated, and cells were cultured for 7 days, 14 days and 21 days in complete medium. After each incubation period, cells were rinsed in PBS (x3) and lysed with 300 µl of M-PER. The number of cells adhered was determined using the *Cytotoxicity Detection Kit* *PLUS (LDH)* as described above.

2.12. Cell differentiation

Lysed cells from the proliferation assay were also used for alkaline phosphatase (ALP) detection using the SensoLyte® pNPP Alkaline Phosphatase Assay Kit (AnaSpec Inc., USA) following the manufacturer's guidelines. Reactions were incubated at 37 ºC for 30 min. After

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incubation, values of absorbance were read in a PowerWave HT microplate reader at 405 nm. A calibration curve was prepared using purified ALP from the kit. Results were normalized versus cell number and time of incubation.

2.13. Statistical analysis

Experiments were performed in duplicate using three replicates per each group. All data are presented as mean values ± standard deviations. A non-parametric Kruskal-Wallis test followed by Mann-Whitney test with Bonferroni correction was used to determine statistical significant (p-value<0.05) differences between the means of the different groups.

3. RESULTS

3.1. Physicochemical characterization

3.1.1 QCM-D

The thickness and surface mass density of the adlayer of immobilized biomolecule on silanized Ti sensors determined by QCM-D analysis is shown in Table 1. Figure S1 (supplementary material) shows the changes in frequency and dissipation over time recorded during the adsorption of the biomolecules. In all cases, the frequency rapidly decreased after injection, reaching values of saturation after 10 and 15 min for the peptides and protein fragments, respectively. Moreover, dissipation coefficients showed small changes during the adsorption of peptides, while very high values were reached for HBII and to a slightly lower extent for the CAS protein fragment. The final adsorbed mass was considerably higher for the protein fragments (609.1 ng/cm^2 for CAS, 823.8 ng/cm^2 for HBII) than for the peptides (220.1 ng/cm^2 for linRGD and 200.0 ng/cm^2 for cRGD).

3.1.2. XPS

The analysis of the chemical composition of the surfaces by XPS is summarized in Table 2. In general, the presence of the biomolecules increased the percentage of C 1s and N 1s and decreased the observed amounts of O 1s, Si 2p and Ti 2p compared to control (Ctrol) non-functionalized surfaces. These effects were more pronounced for the protein fragments (CAS and HBII) than for the peptides (linRGD and cRGD).

3.2. Cell adhesion

3.2.1. Number of cells adhered

The number of cells adhered onto each biofunctionalized surface (Fig. 1) differed depending on the immobilized molecule. Surfaces coated with linRGD, cRGD, CAS or HBII, respectively, showed a significant improvement on cell adhesion compared to FN ($p < 0.05$). There were no statistical differences on cell numbers between the peptides and the protein fragments. On the other hand, the different combinations of CAS and HB II fragments triggered different cell adhesion responses. Low concentrations of CAS (30% and 50%) stimulated an adhesion similar to that of FN, whilst a higher concentration (70%) originated a response comparable to that of linRGD, cRGD, CAS or HBII fragments alone.

3.2.2. Cell spreading

The area of cells adhered on the different biofunctionalized surfaces is shown in Fig. 2. The spreading obtained for cells in contact with linRGD and HBII respectively, was statistically lower ($p < 0.05$) than that obtained for cells in contact with FN, cRGD and CAS. When both CAS and HB II recombinant fragments were mixed, the proportion of each of the two fragments influenced cell spreading. Low percentages (30% and 50%) of CAS induced values of cell spreading close to that found on FN-coated surfaces, whereas higher values of

CAS (70%) led to lower projected cell areas, with values similar to those of the linear RGD peptide and the HBII fragment alone.

3.2.3. Cell morphology

The formation of actin stress fibers (in red) was stimulated by linRGD (Fig. 3B), cRGD (Fig. 3C) and CAS (Fig. 3D), although structures resembling focal contacts were not seen in the cells in contact with these biomolecules. Cells in contact with HBII (Fig. 3E) were not able to generate neither actin filaments nor focal adhesions and showed ruffled borders. Noteworthy, when both CAS and HBII were mixed with low percentages of CAS (Figs. 3F and 3G) cells showed a cytoskeleton conformation closer to that of FN compared to the other biomolecules, whereas the presence of CAS in high percentage (Fig. 3H) showed a cytoskeleton similar to the CAS fragment alone.

3.3. Cell proliferation

The proliferation of rMSCs on the different biofunctionalized substrates after 4 h, 7 days, 14 days and 21 days in culture is shown in Fig. 4. Although cells cultured with linRGD, cRGD, CAS or HBII, respectively, supported cell proliferation within the initial 7 days of incubation, the number of cells on these surfaces did not reach the values obtained with FN coating, which showed the highest values of cell growth at 21 days. Noteworthy, when CAS and HBII recombinant fragments were mixed using less than 70% of CAS, the proliferation levels were closer to FN samples. In contrast, for higher percentages of CAS fragment proliferation rates were similar to short peptides or recombinant fragments separately.

3.4. Cell differentiation

Fig. 5 shows the ALP activity levels of rMSCs after 7, 14 and 21 days in culture. The first 7 days in culture, ALP levels for cells cultured on HBII and high percentage of CAS (70%) were significantly higher compared with FN and the other molecules, which showed levels of

ALP equivalent to that of FN. After two weeks in culture, ALP from cRGD, CAS and CAS (50%) was reduced whilst the other surfaces showed similar levels compared with FN and between them. However, cells cultured on cRGD or CAS fragment exhibited an increase on ALP levels 21 days after seeding, reaching levels similar to those of FN. Noteworthy, HBII-coated surfaces stimulated significantly higher ALP levels than linRGD, cRGD, CAS and FN at all time points, reaching the highest levels of ALP production after 21 days of incubation. Combination of the HBII with a high percentage of CAS (70%) maintained this high ALP expression. In contrast, ALP levels were significantly reduced when HBII was mixed with lower concentrations of CAS (30% and 50%).

4. DISCUSSION

One approach to improve osseointegration in metallic implants is the immobilization of bioactive molecules on their surface. Such modifications are usually achieved by physisorption or covalent bonding. Physisorption is a non-covalent method based on Van der Waals interactions and hydrogen bonds, which are weak and unstable interactions. In contrast, the use of silane chemistry is a preferred and well-extended method for covalently bounding bioactive molecules onto metallic surfaces as it provides a strong and stable binding on the surfaces.[22,47] In this regard, APTES has been extensively used for surface functionalization purposes and demonstrated its suitability for the immobilization of bioactive molecules onto Ti[22] and TiNbHf alloy.[48] Moreover, in a recent study we demonstrated that the use of APTES provides a more homogenous layer of biomolecule attachment in both Ti and TiNbHf alloy compared to physisorption.[49] Hence, in the present work APTES was chosen as linker molecule to immobilize the different bioactive molecules onto a recently developed low modulus TiNbHf alloy.

As previously introduced, the binding of synthetic RGD peptides to Ti surfaces and TiNbHf alloys using this silane has been recently characterized, and yielded a dense and homogenous

layer of peptide on the surfaces.[49] The number and density of molecules immobilized on these surfaces with this method has been reported in a large number of previous studies.[50,51] However, no literature was available on the relative adsorption kinetics of the CAS and HBII fragments on TiNbHf via APTES. QCM-D results indicated a successful attachment of both proteins and peptides on the Ti sensors, reaching values of saturation after 10 min and 15 min of incubation, respectively. Rinsing in PBS resulted in minor protein detachment, proving the stability of the coatings. The amount of adsorbed protein was slightly different for each fragment (i.e. HBII adsorbed mass was 1.3 fold higher than that of CAS). This observation might be attributed to the distinct conformation and chemical sequence of the proteins, which in turn results in different values of protein binding. As expected, surface mass density was lower for peptides compared to protein fragments. The amount of adhered linRGD was slightly higher than cRGD but differences were not statistically significant. Dissipation shift plots indicated that during HBII protein adsorption, water was incorporated and trapped in the layer of protein, as seen in the increasing ΔD. CAS protein layer was less viscoelastic and both peptide adsorbed layers were more rigid compared to the HBII protein fragments layer.

XPS data confirmed QCM-D results and showed a successful biofunctionalization of the surfaces. The observed increases in C 1s, and especially in N 1s, are common indicators of biomolecule attachment (i.e. amide bonds).[22,27,50,51] The reduction in the detectable signals of Ti 2p, O 1s, and Si 2p is also consistent with the presence of biomolecules, which mask the detection of Ti oxide and the silane layer. The binding of the protein fragments yielded the highest variations in chemical composition (e.g. the percentage of N 1s) in comparison to the RGD peptides. This observation is well in agreement with QCM-D values, which indicated that the amount of proteins bound to the surfaces was higher than that of the peptides.

The interaction of cells with the different functionalized surfaces exhibited substantially different behaviours. Both recombinant FN fragments, CAS and HBII, stimulated better cell

adhesion than the whole FN, and similarly to RGDs. The higher activity of a recombinant FNIII$_{7-10}$ fragment compared to plasma human FN has been previously described by the group of García[30] and associated with an enhanced affinity for integrin α5β1. The higher number of attached cells observed for RGD peptides could also be attributed to a higher availability of cell adhesive motifs per area compared to native FN. Although previous studies demonstrated better cell adhesion on CAS fragment than on RGDs,[28–30] these differences tend to disappear in saturated surface densities.[28]   Noteworthy, HBII also stimulated high cell adhesion numbers. This may be explained by the fact that cell surface HSPGs may act as receptors for the HB fragment[52,53] and that integrin receptors α4β1 and αvβ5 may interact with specific sequences in this region of FN.[52,54]

Although the number of cells was high in both FN fragments and RGDs, these cell adhesive motifs were not enough to stimulate cell spreading to similar levels than the native FN. In addition, the number of focal adhesions on these surfaces was lower compared to FN, suggesting that their formation depends on the interaction of cell adhesion sequences with other binding sites present in FN.[55] This scenario completely changed when both recombinant FN fragments were mixed on the same surface. Although the number of cells increased while CAS concentration was incremented, cellular spreading, cytoskeletal organization and focal adhesion formation were improved with higher concentrations of HBII (≥50%). The observed critical biological role of the HBII domain is due to the presence of actin-organizing elements located in the type III$_{13}$ repeat that stimulate the formation of focal adhesions.[55] In this study we have shown that the HBII fragment alone does not support cell spreading and cytoskeletal formation on the surfaces, however, in combination with small percentages of the CAS domain effectively stimulates the formation of stress fibers and focal contacts, reaching cell morphologies similar to FN.[54,56,57]

Integrin mediated adhesion plays a major role in the regulation of cell proliferation, since it stimulates crucial regulatory pathways of cell growth.[58] However, proliferation not only depends on cell adhesive mechanisms but also on the interaction with soluble growth factors. The growth factor receptor signalling is moreover highly regulated by integrin expression and activity.[59,60] In the present study, RGDs and the recombinant FN fragments did not induce cell proliferation at levels comparable to that of full-length FN, probably due to the lack of cooperation between adhesion and growth factor signalling. In contrast, the combination of the CAS with high concentrations of the HBII domain ($\geq$ 50%) effectively improved proliferation of rMSCs. This could be attributed to the fact that besides the cell adhesion cooperative activity, HBII may also act as a growth factor binding domain. This enhancement in cell proliferation was also observed in previous studies by Kim *et al.*[53] and Martino *et al.,*[33] although in their studies a molecule containing both CAS and HBII fragments fused was used instead.

Finally, the stimulation of MSCs to differentiate into the osteoblastic lineage, also known as osseoinduction, is a critical step to ensure the biomaterial osseointegration. The secretion of ALP has been widely used as a marker for osteoblastic differentiation. In the present study, surface functionalization with a synthetic linRGD peptide was enough for stimulating rMSCs differentiation despite the low integrin-specificity described for linear peptides.[36,61]

In the present study, surface functionalization with a synthetic linRGD peptide was enough for stimulating rMSCs differentiation. However, presentation of the RGD motif in a cyclic conformation reduced the expression of ALP levels. This result should not come as a surprise because cRGD shows a higher specificity than linRGD for integrins αvβ3 and αvβ5, but not towards α5β1, which is well known to induce osteogenic differentiation.[62,63] Noteworthy, the presence of HBII yielded the highest levels of ALP activity probably due to the aforementioned capacity to bind diverse growth factors, which may include BMP-2 and

BMP-7 present in the serum.[41] However, the combination of HBII and CAS fragments yielded unexpected results: whereas low percentages of HBII retained high values of ALP expression (CAS-HBII 70:30) higher amounts reduced this activity (CAS-HBII 50:50 and CAS-HBII 30:70) (Fig. 5).

The equimolar presentation of CAS and HBII (50:50) should support similar biological profiles than full length FN. As a matter of fact, comparable cell responses were observed in terms of cell spreading, cytoskeleton formation and proliferation (Fig. 6). In this case, the reduced osteogenic capacity could probably respond to two factors: an insufficient quantity of HBII motifs available for growth factor interaction or the absence of other regulatory sequences present in FN, which may account for the osteogenic activity of the native protein.

It was even more intriguing that an increase in the proportion of HBII (CAS-HBII 30:70) did not stimulate osteoblastic differentiation. We hypothesized that the reduced amount of CAS fragment available for integrin binding would not be enough for generating stable cellular adhesions and focal contacts, thereby engaging part of the HBII fragments in cell adhesion/cytoskeleton organization. This would explain why even if the proportion of CAS is reduced in the mixture CAS-HBII (30:70), the other parameters of cell behavior remain comparable to those of FN (Fig. 6). This "co-adhesive" role for HBII would, in turn, diminish its capacity to sequester growth factors and thus its osteogenic activity. Only when a larger amount of HBII is present (HBII alone) both an adhesive and an osteogenic effect can be maintained.

This hypothesis could also explain the opposite situation (CAS-HBII 70:30). In this context, the high amount of CAS would be sufficient to efficiently stimulate cell adhesion and cytoskeleton organization. This would keep a higher number of HBII fragments free for interaction with growth factors increasing the ALP levels.

5. CONCLUSIONS

In the present study cell response to functionalised Ti25Nb21Hf alloy with two different recombinant FN fragments was studied. The results demonstrate that the mixture of CAS and HBII fragments improves cellular behaviour compared to fragments alone or synthetic RGD peptides, being a promising strategy for biomaterial implantation purposes. This response may be modulated depending on the percentage of each fragment.

When dealing with combinations of molecules that are bifunctional and/or share a similar biological function, it is important to highlight that reducing the percentage of one of the bioactive molecules in the mixture can modify the biological profile of the other one. In the present study, we observed that the reduction on the amount of cell adhesive motifs (CAS) in the mixture was accompanied with a decrease in the osteogenic activity of the HBII fragment. We hypothesized that this effect was attributed to the fact that the low percentage of CAS engaged HBII in cell adhesive processes, thus reducing its capacity to interact with growth factors and hence its osteogenic capacity.

Therefore, it should be taken into account that the equimolar presentation of two bioactive molecules may not always be an optimal approach when these molecules have partially overlapping biological roles. In these instances, the study of different percentages of each motif is recommended.

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TABLES

Table 1. QCM-D study of the thickness and surface mass density of the bound biomolecules to silanized Ti sensors calculated using the Voigt model.

Biomolecules	Thickness (nm)	Surface mass density (ng/cm^2)
CAS	5,54 (±0,09)	609,14 (±10,19)
HBII	7,49 (±0,13)	823,80 (±14,00)
linRGD	2,00 (±0,17)	220,12 (±9,17)
cRGD	1,82 (±0,14)	200,01 (±14,98)

Table 2. Surface chemical composition (expressed as atomic percentages) of the biofunctionalized samples by XPS.

	C 1s	N 1s	O 1s	Si 2p	Ti 2p
Ctrol	22,96 (± 1.87)	0,76 (± 0.19)	56,25 (± 1.45)	0,49 (± 0.08)	19,57 (± 0.15)
CAS	47,63 (± 0.13)	10,57 (± 0.25)	31,93 (± 0.04)	3,01 (± 0.06)	6,87 (± 0.23)
HBII	46,45 (± 1.48)	10,26 (± 0.53)	32,85 (± 1.22)	3,66 (± 0.08)	6,78 (± 0.72)
linRGD	32,16 (± 0.42)	4,82 (± 0.03)	46,31 (± 0.33)	4,82 (± 0.33)	11,91 (± 0.28)
cRGD	34,77 (± 1.07)	5,33 (± 0.05)	43,55 (± 0.76)	5,53 (± 0.04)	10,83 (± 0.30)

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FIGURE CAPTIONS

Fig. 1. Adhesion of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules after 4 h of culture. Letter "a" indicates study groups with no statistical differences with FN. Letter "b" indicates study groups with statistically significant differences compared to FN.
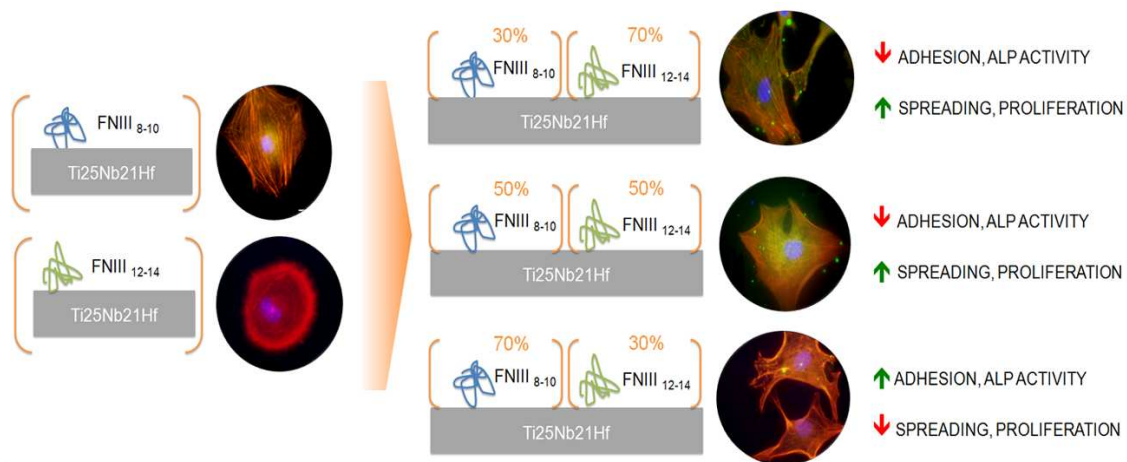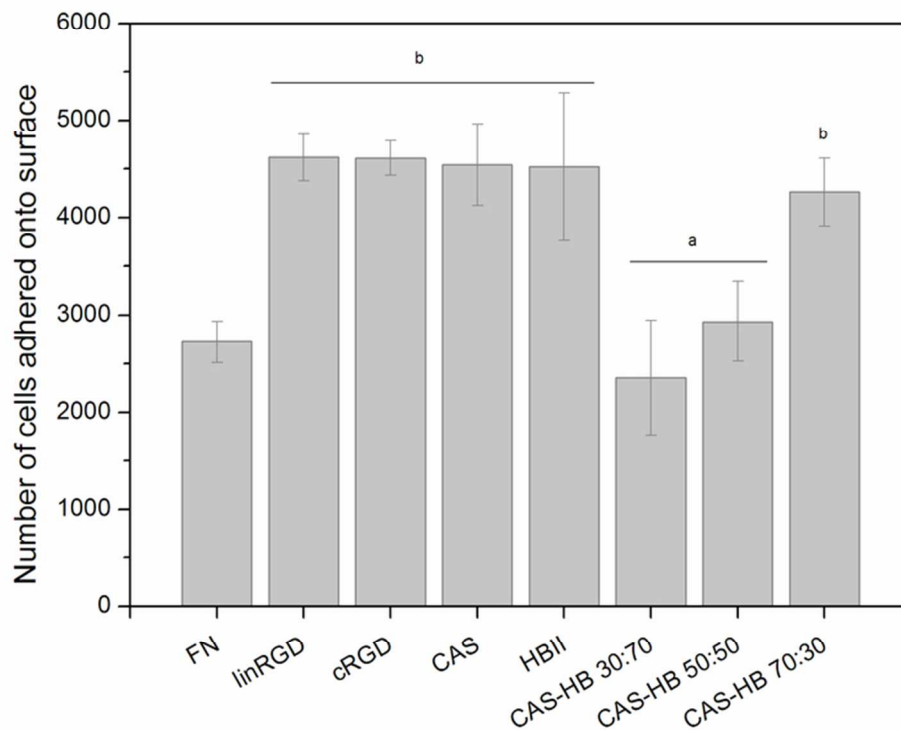
Fig. 2. Spreading of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules after 4 h of culture. Letter "a" indicates study groups with no statistical differences with FN. Letter "b" indicates study groups with statistically significant differences compared to FN.

Fig. 3. Representative high magnification images (60x) of rMSCs spreading and morphology after 4h in culture with FN (A), linRGD (B), cRGD (C), CAS (D), HB II (E), CAS-HBII 30:70 (F), CAS-HBII 50:50 (G) and CAS-HBII (H) biofunctionalized surfaces. Scale bar 20 μm.

Fig. 4. Proliferation of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules. At each time, "a", "c", "g" and "k" indicates no statistical differences compared to FN while distinct letters denote statistical differences to FN and between conditions.

Fig. 5. ALP activity of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules. At each time, "a", "c" and "e" indicates no statistical differences compared to FN while distinct letters denote statistical differences to FN and between conditions.

Fig. 6. Summary of cell responses to functionalized surfaces with FN, different mixtures of CAS and HBII fragments (30:70, 50:50 and 70:30) and CAS and HBII fragments alone. Biomolecules with similar responses were grouped by colors for better interpretation of results.

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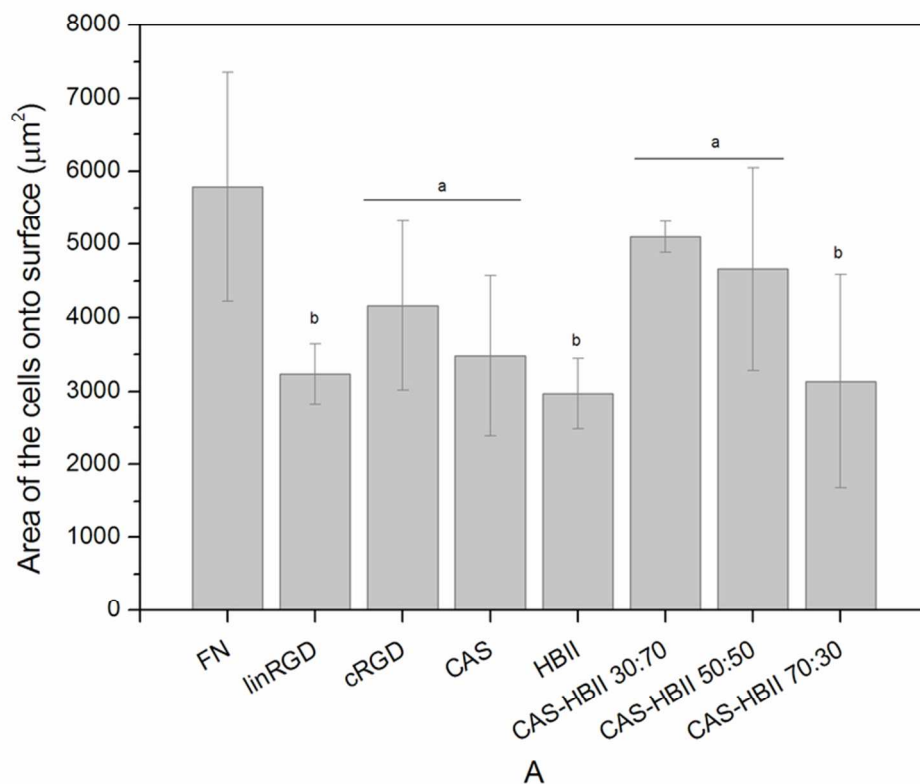42
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Fig. 1. Adhesion of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules after 4 h of culture. Letter "a" indicates study groups with no statistical differences with FN. Letter "b" indicates study groups with statistically significant differences compared to FN.
68x55mm (300 x 300 DPI)

Fig. 2. Spreading of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules after 4 h of culture. Letter "a" indicates study groups with no statistical differences with FN. Letter "b" indicates study groups with statistically significant differences compared to FN.
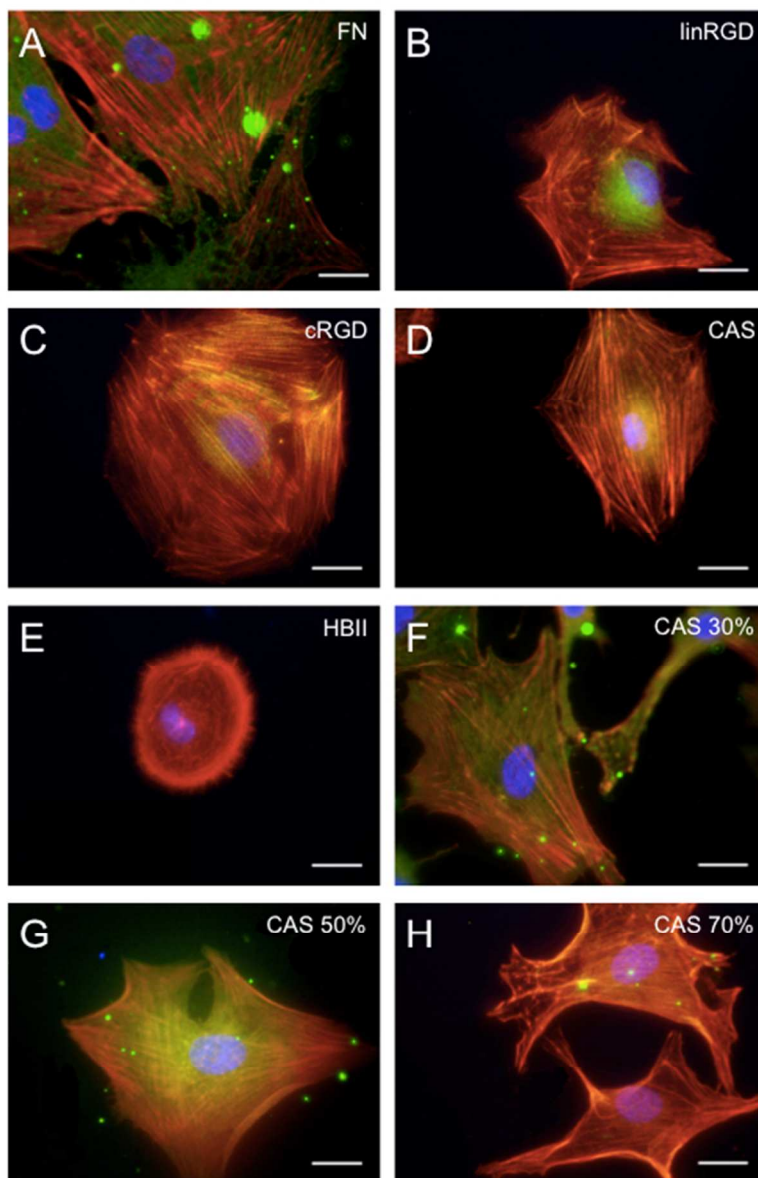73x63mm (300 x 300 DPI)

Fig. 3. Representative high magnification images (60x) of rMSCs spreading and morphology after 4h in culture with FN (A), linRGD (B), cRGD (C), CAS (D), HB II (E), CAS-HBII 30:70 (F), CAS-HBII 50:50 (G) and CAS-HBII (H) biofunctionalized surfaces. Scale bar 20 µm.
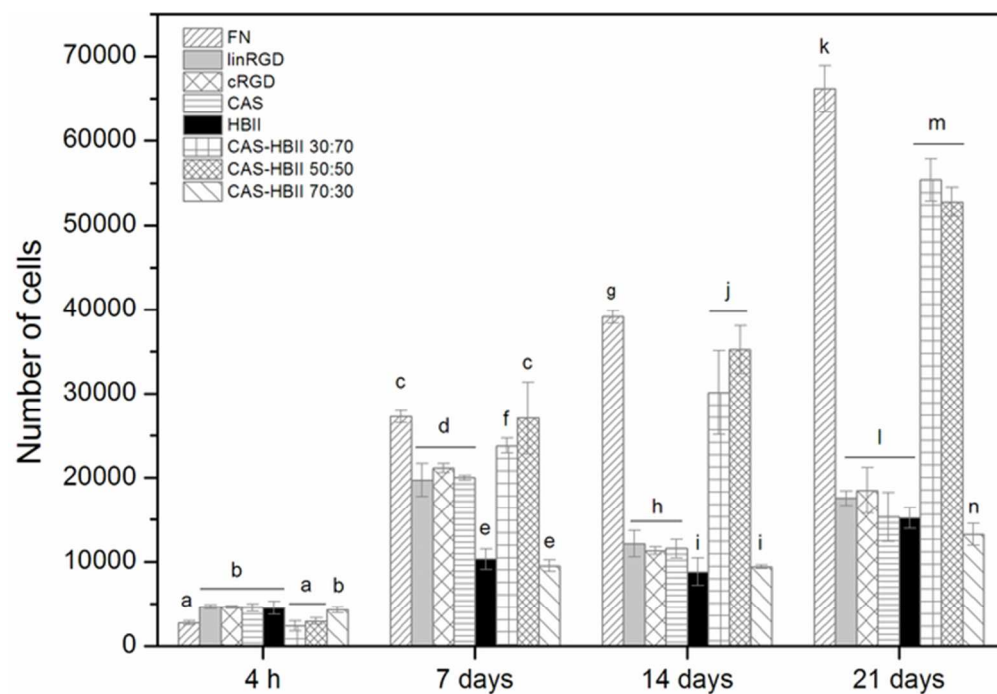42x65mm (300 x 300 DPI)

Fig. 4. Proliferation of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules. At each time, "a", "c", "g" and "k" indicates no statistical differences compared to FN while distinct letters denote statistical differences to FN and between conditions.
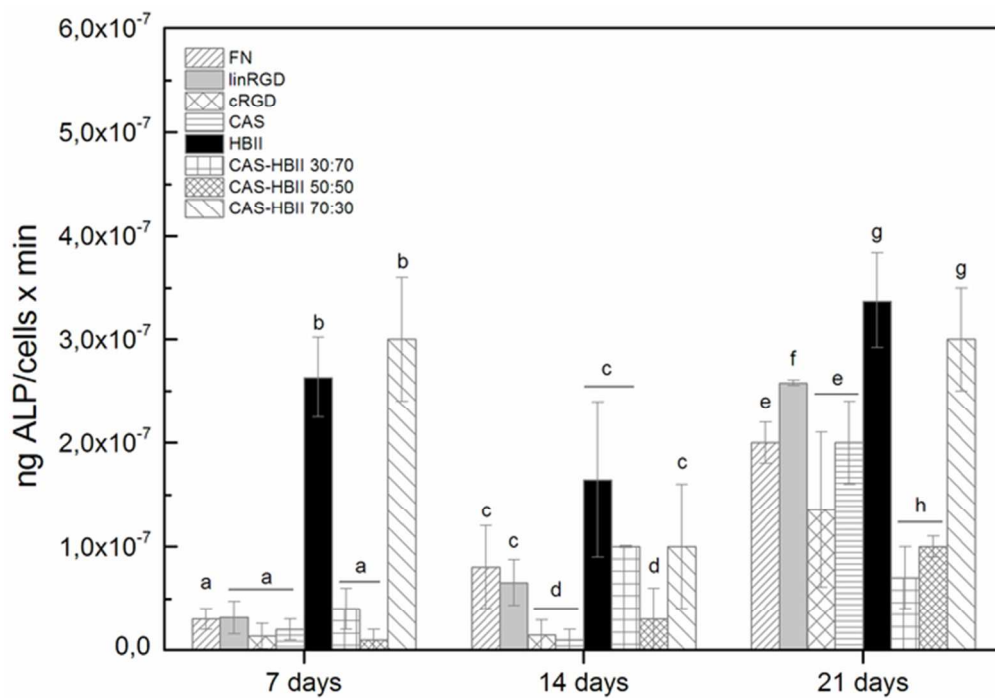58x40mm (300 x 300 DPI)

Fig. 5. ALP activity of rMSCs onto Ti25Nb21Hf alloy biofunctionalized with the different biomolecules. At each time, "a", "c" and "e" indicates no statistical differences compared to FN while distinct letters denote statistical differences to FN and between conditions.
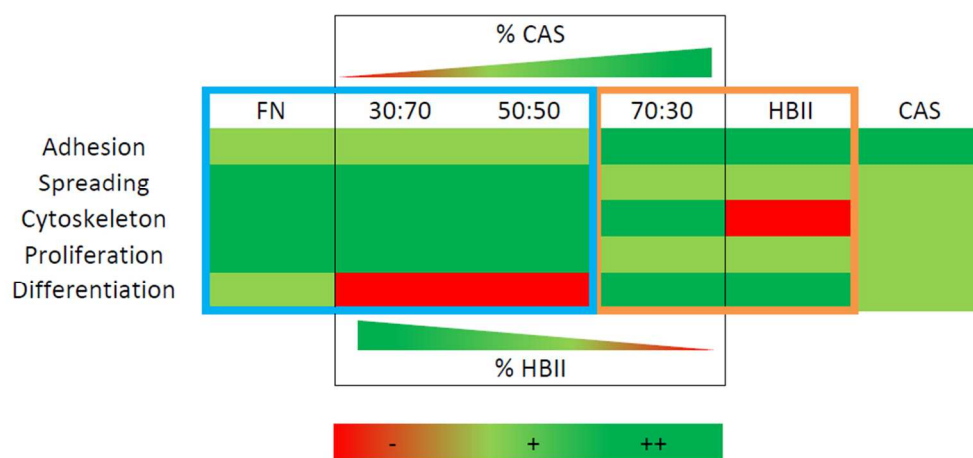59x42mm (300 x 300 DPI)

Fig. 6. Summary of cell responses to functionalized surfaces with FN, different mixtures of CAS and HBII fragments (30:70, 50:50 and 70:30) and CAS and HBII fragments alone. Biomolecules with similar responses were grouped by colors for better interpretation of results.
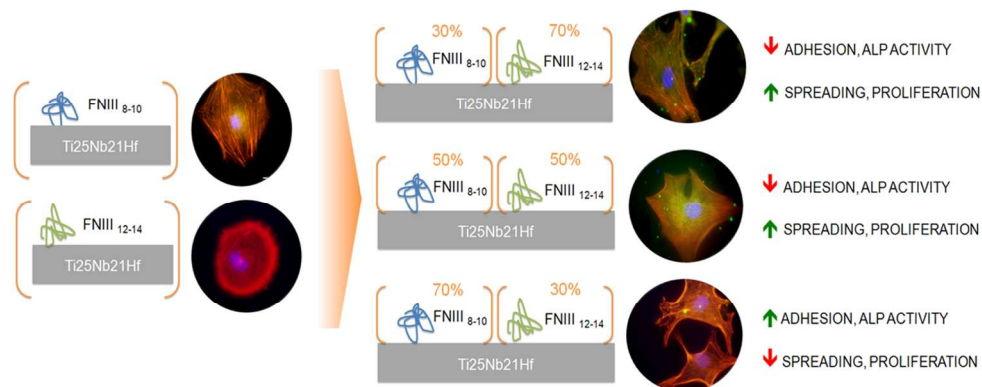84x40mm (300 x 300 DPI)

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112x44mm (300 x 300 DPI)