

DOI: 10.1002/((please add manuscript number))

**Article type: Communication**

**Title: Enhanced biological activity of BMP-2 bound to surface-grafted heparan sulfate**

*Elisa Migliorini\*, Patrick Horn, Tamás Haraszti, Seraphine V. Wegner, Christian Hiepen, Petra Knaus, Ralf P. Richter and E. Ada Cavalcanti-Adam\**

Dr. E. Migliorini, Dr. S. V. Wegner, Prof. E.A. Cavalcanti-Adam  
Department of Biophysical Chemistry, Institute of Physical Chemistry, Heidelberg University,  
Im Neuenheimer Feld 253, 69120 Heidelberg, Germany  
Department of Cellular Biophysics, Max Planck Institute for Medical Research, Postal  
Address: Heisenbergstr. 3 D-70569 Stuttgart, Germany  
E-Mail: [migliorini@is.mpg.de](mailto:migliorini@is.mpg.de); [ada.cavalcanti-adam@urz.uni-heidelberg.de](mailto:ada.cavalcanti-adam@urz.uni-heidelberg.de)

Dr. P. Horn

Department of Medicine V, Heidelberg University, INF 410, 69120 Heidelberg, Germany

Dr. T. Haraszti

DWI - Leibniz Institute for Interactive Materials, Forckenbeckstr. 50 52056 Aachen, Germany

Dr. S.V. Wegner

Department of Biophysical Chemistry, Institute of Physical Chemistry, Heidelberg University,  
Im Neuenheimer Feld 253, 69120 Heidelberg, Germany

Max Planck Institute for Polymer Research, Ackermannweg 10, 55128 Mainz, Germany

Dr. C. Hiepen, Prof. P. Knaus

Institute of Biochemistry, Freie Universität Berlin, Thielallee 63, 14195 Berlin, Germany

Dr. R. P. Richter

University of Leeds, School of Biomedical Sciences and School of Physics and Astronomy,  
Leeds, LS2 9JT, UK; CIC biomaGUNE, Biosurfaces Lab, Paseo Miramon 182, 20009 San  
Sebastian, Spain; University Grenoble-Alpes and CNRS, Laboratory of Interdisciplinary  
Physics, 140 Rue de la Physique, 38402 St. Martin d'Hères, France

Keywords: BMP-2, heparan sulfate, extracellular matrix, noggin, biomimetic platforms

Bone morphogenetic protein-2 (BMP-2) is known to initiate the differentiation of mesenchymal stem cells (MSCs) into osteoblasts and chondrocytes *in vivo* and *in vitro*,<sup>[1]</sup> as well as the transdifferentiation of mouse myoblasts C2C12 cell line into mineralizing bone-like cells.<sup>[2, 3]</sup> The BMP-2 signaling pathway is activated by the binding of the growth factor to two types of transmembrane serine/threonine kinase receptors, the BMP type I (BMPRI) and type II (BMPRII) receptors. Binding of BMP-2 to BMPRI and BMPRII induces the phosphorylation of the SMAD1/5/8 complex, which then together with co-SMAD (SMAD4) translocates to the nucleus and, with other DNA-binding proteins, participates in the transcriptional regulation of osteogenic target genes.<sup>[4]</sup>

1 The clinical use of BMP-2 was approved in 2002 by the Food and Drug Administration in the  
2 USA and validated by the national Medical Agencies in Europe. BMP-2 has been rapidly  
3 introduced into orthopedic clinical practice<sup>[5]</sup> but, in some cases, even high doses (up to 2  
4 mg/level) appear only marginally effective, likely due to an autocrine BMP-2 inhibition by  
5 noggin the major extracellular BMP antagonist, at sites of BMP-2 application.<sup>[6]</sup> Therefore,  
6 improving BMP-2 activity, by using delivery systems that suppress BMP-2 inhibition by  
7 noggin, would help in overcoming current challenges in BMP-2 clinical application.<sup>[7]</sup>

8 The development of materials that are able to control BMP-2 molecular presentation and local  
9 concentration, is an essential approach for a deeper understanding of BMP-2 functions and the  
10 modulation of its biological activity.<sup>[8]</sup>

11 In tissues, BMP-2 is bound to extracellular matrix (ECM) components such as proteins<sup>[9]</sup> and  
12 glycosaminoglycans (GAGs), in particular to heparan sulfate (HS)<sup>[10, 11]</sup>. BMP-2 is a  
13 homodimeric protein and has been reported to bind *in vitro* to heparin (Hp), a highly sulfated  
14 form of HS, *via* a binding site located at the N-terminus (**Figure 1A**) with an affinity of  
15 approximately 20 nM.<sup>[12]</sup> The binding of BMP-2 to HS, on the other hand, remains to be fully  
16 characterized in terms of stoichiometry, affinity and concomitant HS morphological changes.  
17 HS, a linear polysaccharide formed by variably sulfated repeating disaccharide units, is  
18 typically covalently attached to core-proteins forming HS proteoglycans (HSPGs). The  
19 functions of HSPGs depend on their structure and location, being either at the cell surface or  
20 as part of the ECM.<sup>[13]</sup>

21 Contradictory effects of cell surface HSPGs on BMP-2 activity have been reported. On the  
22 one hand, endogenous HSPGs negatively modulate both chondrogenic<sup>[14]</sup> and osteogenic<sup>[10]</sup>  
23 differentiation by inhibiting the activation of BMP-2 signaling. On the other hand, HSPGs act  
24 as BMP-2 co-receptors, promoting the formation of complexes between BMPRII and BMPRI,  
25 thereby enhancing the bioactivity of BMP-2.<sup>[15]</sup> Interestingly, the BMP antagonist noggin also  
26 binds to HS.<sup>[16]</sup> Once bound to cell surface HSPGs, noggin maintains its function to inhibit

1 the activity of BMP-4, another member of the BMP family, by blocking the binding epitopes  
2 for BMPRI and BMPRII.<sup>[16, 17]</sup>  
3

4 To date, the role of HSPGs in the extracellular space (ECM-HSPGs) on BMP-2 bioactivity is  
5 only partially explored. The main functions so far attributed to ECM-HSPGs are to prevent  
6 the diffusion of growth factors away from the regions where they are likely to be required.<sup>[18]</sup>  
7  
8 However, it is still unknown if these HSPGs can positively modulate BMP-2 biological  
9 activity thus enhancing *in vitro* osteogenic differentiation.<sup>[19]</sup>  
10  
11

12 The presence of HS added to the culture media of BMP-responsive cells (C2C12 myoblasts)  
13 prolongs SMAD 1/5/8 phosphorylation in these cells and reduces the interactions with the  
14 antagonist noggin,<sup>[11, 20]</sup> in contrast to the effect of cell surface HSPGs.<sup>[16]</sup> Since *in vivo* HS is  
15 not in solution, but rather covalently bound to core-proteins through its reducing end forming  
16 HSPGs,<sup>[21]</sup> administering HS in the cell culture media is unlikely to be representative of the *in*  
17 *vivo* extracellular HS.  
18  
19

20 Biomimetic platforms presenting HS in a bound and oriented manner are therefore suitable to  
21 explore (i) the characteristics of HS/BMP-2 binding at the molecular and cellular levels, (ii)  
22 the impact of BMP-2 presentation on BMP-2 activity, and (iii) the interaction of noggin with  
23 grafted HS and its recognition of the HS-bound BMP-2. The strength of such platforms lies in  
24 the precise molecular presentation, which gives the ability to control and characterize BMP-2  
25 interactions with HS, as well as the possibility to further exploit them as substrates for  
26 investigations on the impact of HS on BMP-2 activity in cells. We designed our model  
27 surfaces in a way that HS is grafted by its reducing end,<sup>[22]</sup> mimicking its attachment to HSPG  
28 core proteins *in vivo*. To this end, we functionalized the reducing end of HS with biotin and  
29 used gold-coated surfaces functionalized first with a mixed monolayer of oligoethyleneglycol  
30 (OEG)-thiol and biotin-OEG-thiol and then with a streptavidin monolayer as substrates<sup>[23]</sup>  
31  
32  
33  
34  
35  
36  
37  
38  
39  
40  
41  
42  
43  
44  
45  
46  
47  
48  
49  
50  
51  
52  
53  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65  
**(Figure 1B)**. The OEG monolayer passivates against non-specific binding of proteins, here  
tested using bovine serum albumin (BSA) **(Figure S1)**, whereas streptavidin (SAv) binds

1 through a specific interaction with the biotin end-groups and forms a rigid monolayer. Such  
2 monolayer serves then as a mediator for the immobilization of biotinylated compounds such  
3  
4 as biotinylated HS (b-HS; **Figure 1C**) and biotinylated BMP-2 (b-BMP2; **Figure 1D**). BMP-  
5  
6 2 binds to SA<sub>v</sub> *via* a biotin-PEG<sub>12</sub>-NHS ester linker (of ~5.6 nm contour length) which reacts  
7  
8 with primary amines of the growth factor. Surface functionalization steps, specificity of the  
9  
10 molecular binding and mechanical properties of the biomolecular films were characterized by  
11  
12 quartz crystal microbalance with dissipation monitoring (QCM-D); adsorption and desorption  
13  
14 rates of the different biomolecules, as well as biomolecular surface densities, were quantified  
15  
16  
17 by spectroscopic ellipsometry (SE).  
18  
19  
20

21 **Figure 1C** and **D** shows QCM-D characterization of the two biomimetic platforms ( $\Delta f$  and  
22  
23  $\Delta D$  correspond to the fifth overtone). After the formation of a SA<sub>v</sub> monolayer with an  
24  
25 expected thickness of approximately 4 nm<sup>[24]</sup> (determined using Sauerbrey's equation;  
26  
27 **Figure 1C**, 60 to 80 min), b-HS adsorbs rapidly forming a hydrated and soft film, indicated  
28  
29 by the increase in dissipation of  $4.5 \pm 0.5 \times 10^{-6}$  (**Figure 1C**, 90 to 110 min). BMP-2,  
30  
31 incubated at a concentration of 96 nM, stably binds to the b-HS film causing an increase in  
32  
33 film rigidity, as demonstrated by the negative dissipation shift of  $-0.9 \pm 0.1 \times 10^{-6}$  (**Figure 1C**,  
34  
35 150 to 180 min). Complementary assays based on fluorescence recovery after photobleaching  
36  
37 (FRAP), using films of b-HS grafted to supported lipid bilayers in the fluid phase, revealed  
38  
39 that BMP-2 binding reduces the lateral mobility of HS chains substantially. This suggests that  
40  
41 the increased rigidity arises from cross-linking of HS chains mediated by the growth factor  
42  
43 (**Figure S2**). This phenomenon has been reported for other ECM signaling proteins, such as  
44  
45 chemokines and growth factors, some of which present multiple HS binding sites.<sup>[26]</sup> We  
46  
47 speculate that HS cross-linking induced by BMP-2 might be due to the presence of two or  
48  
49 more independent Hp/HS binding sites at the N-terminus region of the BMP-2 dimer, as  
50  
51 previously predicted.<sup>[25]</sup> A mutant form of BMP-2, EHBMP-2, where the N-terminal region  
52  
53 responsible for the Hp binding was substituted by a heterologous sequence from human  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65

interleukin-2,<sup>[12]</sup> does not bind to b-HS (**Figure 1C**, grey curve) and does not induce HS film cross-linking (**Figure S2**). This demonstrates that BMP-2 binds HS specifically through the same site as Hp.

For comparison, we also adopted a second BMP-2 immobilization strategy that does not involve HS. In this case, biotinylated BMP-2 (b-BMP2) binds directly to the SAV monolayer until saturation (**Figure 1D**, 50 to 110 min). When incubated at the same concentration (96 nM) without the biotin tag, BMP-2 does not bind stably (**Figure 1D**, gray curve) demonstrating the b-BMP2 is specifically attached *via* biotin to the SAV monolayer.

To quantify the binding strength of BMP-2 to b-HS films, we used spectroscopic ellipsometry (SE) and performed a titration assay (**Figure S3A**). The titration curve (**Figure S3B**) was approximated well by the simple Langmuir isotherm:

$$\Gamma_{eq} = \Gamma_{max} \frac{[BMP-2]}{K_d + [BMP-2]}, \quad (1)$$

with an affinity of  $K_d \approx 1.6 \mu\text{M}$  and a maximal BMP-2 surface density of  $\Gamma_{max} \approx 1000 \text{ ng/cm}^2$ .

For this analysis, it has to be considered that HS is not a homogeneous polymer: the constituent monosaccharides are variably sulfated and/or might exist as different epimers, (regions of high sulfation coexisting with regions of low sulfation along individual HS chains).

These structural features can also vary from one HS source to another.<sup>[27, 28]</sup> It is therefore likely that HS presents a spectrum of binding sites rather than a single type. The affinity constant of  $1.6 \mu\text{M}$  should thus be considered as an effective value, resulting from a spectrum of binding sites with different affinities.<sup>[29]</sup> For comparison, an affinity of 20 nM for Hp has been previously reported,<sup>[12]</sup> indicating that a high degree of sulfation might indeed substantially enhance the effective binding strength in comparison to the value obtained here.

At the maximal surface density predicted by the Langmuir isotherm ( $\Gamma_{max} = 1000 \text{ ng/cm}^2$ ), we calculated that up to 11 BMP-2 dimers would bind to a 12 kDa b-HS chain; in such condition ~2.1 HS disaccharides are available on average per BMP-2 dimer. A small non-specific

binding of BMP-2 to the SAV monolayer has to be considered when high BMP-2 concentrations are used (**Figure S4A**). At high concentration it is also possible that several BMP-2 molecules interact with each other forming aggregates (**Figure S4B**). Moreover, BMP-2 in solution has been found to have a limited physical stability, with aggregates of various sizes forming in a pH-dependent manner.<sup>[30]</sup> Future studies using shorter oligosaccharides (down to 3 disaccharides) would be useful to study the minimal HS/Hp disaccharides length able to bind BMP-2.

The desorption of BMP-2 upon buffer rinsing is well described by the exponential function (**Figure S3A**, red line):

$$\Gamma = \Gamma_{\text{r}} e^{-k_{\text{off}} \Delta t} + \Gamma_{\text{ir}} \quad (2)$$

with an apparent off-rate  $k_{\text{off}} = 6.1 \pm 1.9 \times 10^{-4} \text{ s}^{-1}$ .  $\Gamma_{\text{ir}}$  and  $\Gamma_{\text{r}}$  correspond, respectively, to BMP-2 fractions that are irreversibly and reversibly bound to the b-HS film. The fit reveals similar values for  $\Gamma_{\text{ir}}$  and  $\Gamma_{\text{r}}$ , meaning that approximately 50% of BMP-2 is released from the b-HS film. The heterogeneous structure of HS chains<sup>[28]</sup> and the spectrum of binding sites and affinities that results from it may well explain the presence of a BMP-2 fraction that binds HS reversibly and another fraction that binds HS stably.

SE was also used to control the surface density of the active biomolecules, a fundamental piece of information to perform studies on BMP-mediated cellular responses (**Figure 2**). To this end, the assembly of biomimetic surfaces was followed step by step, as for the QCM-D measurements. The areal mass densities obtained from the SE data are reported in **Table 1**. b-HS binds to SAV (**Figure 2A** and **Table 1**) and, considering SAV molecular mass of 60 kDa, the amount of HS bound on average per available biotin-binding site (assuming that two of four sites engage in the immobilization to the surface) is  $6 \pm 1.6 \text{ kDa}$ . This value is below the average HS molecular mass employed (12 kDa). As previously discussed<sup>[23]</sup>, this discrepancy is likely to be due to the large size distribution of HS in solution, i.e. capture on SAV has selected the shorter chains in the initial HS sample.

1 Thanks to the quantification of surface densities afforded by SE, we estimate that, at 96 nM  
2 concentration, each BMP-2 dimer has approximately 20 kDa of HS, corresponding to roughly  
3  
4 36 disaccharides, available on average at signal stabilization.  
5

6  
7 Noggin, does not bind to SAV (**Figure S5A**), as expected, but has a HS/Hp binding site<sup>[31]</sup> and  
8  
9 indeed readily binds to the native b-HS film, at an areal mass density of  $\sim 112 \pm 0.7$  ng/cm<sup>2</sup>  
10  
11 when incubated at  $\sim 200$  nM (**Figure S5B**). When BMP-2 loaded b-HS film is incubated with  
12  
13 noggin at the same concentration, noggin binds to the film (**Figure 2A** and **Table 1**). The total  
14  
15 amount of b-HS bound proteins (BMP-2 + noggin) is higher than the amount the individual  
16  
17 proteins adsorbed on b-HS. This indicates that both proteins can be present in the b-HS film  
18  
19 simultaneously, however it remains unclear whether b-HS-bound noggin recognizes at the  
20  
21 same time BMP-2.  
22  
23  
24  
25

26  
27 At a concentration of 96 nM, one b-BMP2 binds on average to two SAV molecules (**Figure**  
28  
29 **2B**), which is expected to be due to the larger dimensions of BMP-2 compared to SAV ( $\sim 8$  nm  
30  
31 vs.  $\sim 5$  nm). After immobilization of b-BMP-2 on SAV, we added noggin and observed that it  
32  
33 binds at a ratio of 1 b-BMP2 to 0.7 noggin molecules (**Figure 2A**). As noggin is known to  
34  
35 bind BMPs with a 1:1 stoichiometry,<sup>[17]</sup> this implies that  $\sim 30\%$  of immobilized b-BMP2 is  
36  
37 apparently not recognized by noggin. Plausible explanations are that the noggin binding site  
38  
39 of some BMP-2 molecules is oriented towards the surface and therefore not accessible, and/or  
40  
41 that the biotin-PEG-NHS is reacting with Lys 97 and/or Lys 101, thus protecting the binding  
42  
43 epitope recognised by noggin<sup>[7]</sup> (**Figure 1A**). Noggin inhibits BMP signalling by blocking  
44  
45 BMP binding epitopes for both BMPRI and BMPRII.<sup>[17]</sup> It is therefore likely that all b-BMP2  
46  
47 molecules recognized by noggin are also accessible to cell surface receptors.  
48  
49  
50  
51  
52

53  
54 To understand how the type of BMP-2 immobilization impacts on BMP-2 and noggin  
55  
56 biological activity, we next performed functional experiments using BMP-responsive cells.  
57  
58 Murine C2C12 myoblasts and mesenchymal stem cells from human bone marrow (hMSC)  
59  
60  
61  
62  
63  
64  
65

1 were plated for short periods (30 to 180 minutes) on the biomimetic platforms to compare the  
2 bioactivity of soluble BMP-2 (sBMP-2), immobilized b-BMP-2 and b-HS/BMP-2.  
3

4 We first characterized the signaling response in C2C12 cells, which form myotubes upon  
5 reaching confluency by switching to low serum condition,<sup>[32]</sup> but in presence of BMP-2 their  
6 myogenic differentiation is inhibited resulting in transdifferentiation towards the osteogenic  
7 lineage.<sup>[33]</sup> The phosphorylation level of SMAD 1/5 proteins, which are direct downstream  
8 effectors of the canonical BMP-SMAD signaling pathway, was used as indicator for BMP-2  
9 bioactivity. We previously demonstrated that the covalent immobilization of BMP-2 on a  
10 surface *via* a heterobifunctional chemical linker retained the growth factor's biological  
11 activity and triggered BMP-mediated signaling in C2C12 cells.<sup>[34]</sup> In the present study, we  
12 investigate the effect of BMP-2 surface presentation *via* b-HS, which resembles its  
13 presentation in the ECM, in comparison to its surface immobilization *via* biotin. In both  
14 platforms BMP-2 surface concentration was determined to be in the range of 35 to 50 ng/cm<sup>2</sup>  
15 (**Table 1**). As reference, we used sBMP-2 at a concentration of 20 nM.<sup>[2]</sup> Surfaces presenting  
16 only b-HS were used as negative control. p-SMAD 1/5 phosphorylation kinetics were  
17 determined at 30, 90 and 180 minutes after cell plating (**Figure 3A and C**). While SMAD 1/5  
18 phosphorylation induced by sBMP-2 decreases during a 3-hour stimulation period, the same  
19 could not be observed when BMP-2 is bound to b-HS or in the case of b-BMP2 immobilized  
20 on SAV. In particular, for the latter, SMAD 1/5 phosphorylation peak is delayed at 90 minutes  
21 and remained stable also for 180 minutes, in line with what has been previously observed for  
22 covalently immobilized BMP-2.<sup>[34]</sup> After 180 minutes, the levels of phosphorylated SMAD  
23 1/5 are significantly higher in cells exposed to BMP-2 bound to b-HS than in presence of  
24 sBMP-2. Interestingly, for all the measured time points, a significant enhancement of p-  
25 SMAD 1/5 levels is observed when BMP-2 is presented through b-HS in comparison to b-  
26 BMP-2 (**Figure 3A**). To discard the possibility that b-HS enhances SMAD phosphorylation  
27 independently of sBMP-2, we used the mutated form of BMP-2 unable to bind HS (EHBMP-  
28  
29  
30  
31  
32  
33  
34  
35  
36  
37  
38  
39  
40  
41  
42  
43  
44  
45  
46  
47  
48  
49  
50  
51  
52  
53  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65



1 2) on b-HS platforms. In this setting SMAD 1/5 phosphorylation levels at 180 minutes were  
2 similar for both sBMP-2 and sEHBMP-2, and in the case of EHBMP-2 they were also not  
3 increased by the presence of b-HS (**Figure S6**), suggesting that the enhancement of BMP-2  
4 bioactivity is due to the specific presentation of BMP-2 by b-HS. It is thus conceivable that b-  
5 HS presents BMP-2 in the correct orientation to the BMP receptor complex.  
6

7  
8  
9  
10  
11 Taken together, these results show that both immobilization strategies (b-BMP2 and b-  
12 HS/BMP-2) prolong the biological activity of the growth factor in comparison to its  
13 presentation to cells when added in the culture media. On b-HS presenting surfaces, the  
14 retention of BMP-2 might be favored by the cross-linking of the b-HS film (**Figure S2**). We  
15 further demonstrate that the presentation *via* b-HS enhances BMP-mediated signaling in  
16 C2C12 cells in comparison to its direct immobilization on SA<sub>v</sub>. To test the role of b-HS as  
17 co-factor and to better elucidate the nature of the binding between b-HS and BMP-2, the  
18 affinity of the complex b-HS/BMP-2 to BMP receptors and the functional blocking of HSPGs  
19 should be addressed in future studies.  
20  
21

22 We observed a similar result in the kinetics of SMAD 1/5 phosphorylation in primary hMSCs  
23 in response to the same immobilization strategy (**Figure 3C**). Indeed, on b-BMP2, SMAD 1/5  
24 phosphorylation is prolonged for 180 min, while it decreased over time in presence of sBMP-  
25 2. In contrast to C2C12 cells, the presence of b-HS does not enhance BMP-2 signaling in  
26 hMSCs in comparison to the presentation of BMP-2 immobilized on SA<sub>v</sub>, which generates  
27 also high and sustained levels of p-SMAD 1/5 expression. Furthermore, a short-time  
28 stimulation of hMSCs on b-HS/BMP-2 and b-BMP2 presenting surfaces is sufficient to  
29 promote osteogenic differentiation. Cells plated for 90 minutes on the biomimetic platforms in  
30 the presence of BMP-2, either immobilized or added to the cell media, and then re-plated for  
31 24 days on untreated tissue culture plates, express alkaline phosphatase (ALP), a marker of  
32 osteogenic differentiation,<sup>[1]</sup> without and additional factors (**Figure S7**). The observed ALP  
33 expression in cells exposed to surfaces presenting b-HS and BMP-2 is significantly higher  
34  
35  
36  
37  
38  
39  
40  
41  
42  
43  
44  
45  
46  
47  
48  
49  
50  
51  
52  
53  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65

1 than the expression in cells exposed to sBMP-2. We conclude that early events triggered by  
2 the presentation of BMP-2 via b-HS are sufficient to activate the SMAD1/5 downstream  
3 signaling, which induces hMSCs differentiation towards the osteogenic lineage.  
4

5  
6  
7 To assess whether the presence of b-HS prevents BMP-2 recognition by its antagonist noggin,  
8 as previously suggested for soluble HS (sHS),<sup>[20]</sup> we analyze short-term BMP-2 signaling on  
9 surfaces presenting BMP-2 bound to b-HS or b-BMP2 bound to SA<sub>v</sub> in the presence of a  
10 double molar excess of noggin in solution (**Figure 3B and D**). As comparison, the same molar  
11 ratio has been used for sBMP-2 and for sBMP-2 bound to sHS. C2C12 and hMSCs responded  
12 in a comparable manner, demonstrating that noggin, by occupying to BMPRI and BMPRII  
13 binding epitopes, is able to inhibit the bioactivity of sBMP-2, sBMP-2 bound to sHS and b-  
14 BMP2 grafted on SA<sub>v</sub>. Surprisingly, when BMP-2 is bound to b-HS, the effect of noggin is  
15 negligible. To rule out the effect of BMP-2 released from b-HS (**Figure S3**) we exposed cells  
16 to an excess of soluble noggin, which we expect to antagonize the biological activity of the  
17 released BMP-2. We demonstrate that even in presence of an excess of noggin in solution (ref.  
18 b-HS BMP-2 noggin excess), Smad 1/5 phosphorylation levels remain high.  
19

20  
21 We speculate therefore that noggin does not efficiently recognize BMP-2 bound to b-HS, thus  
22 leaving a sufficient amount of BMP-2 which is bound to b-HS and still accessible to BMPRI  
23 and BMPRII to activate the SMAD 1/5 pathway. Future studies on the competition between  
24 b-HS/BMP-2 complex, noggin and BMP receptors might better clarify the impact of HS on  
25 BMP-2 interaction with noggin and with BMP receptors. Moreover, to consolidate our  
26 observations, platforms presenting controlled HS sulfation patterns and HS from different  
27 sources, are in the scope of future studies.  
28  
29  
30  
31  
32  
33  
34  
35

## 36 CONCLUSIONS:

37  
38 We designed a biomimetic and versatile platform for molecular and cellular studies, which  
39 presents immobilized BMP-2 alone (b-BMP2) or bound to b-HS grafted *via* its reducing end  
40  
41  
42  
43  
44  
45  
46  
47  
48  
49  
50  
51  
52  
53  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65

1 to SA<sub>v</sub> monolayers, similar to BMP-2 presentation by HSPGs in the ECM. By controlling the  
2 surface density and the stoichiometry of all components, we defined the apparent binding  
3 affinity between BMP-2 and b-HS and we demonstrated that BMP-2 can cross-link b-HS  
4 chains likely due to several independent Hp/HS binding sites at the N-terminus. These  
5 platforms represent therefore a versatile and tuneable biomimetic tool able to be exploited as  
6 active substrates for C2C12 and hMSCs stimulation towards osteogenic differentiation. We  
7 show that (i) surface immobilization of BMP-2 prolongs the p-SMAD 1/5 signalling  
8 activation with respect to the soluble BMP-2, (ii) the specific presentation *via* b-HS enhances  
9 the p-SMAD 1/5 levels on C2C12 cells and prevents the antagonistic effect of noggin on both  
10 C2C12 and hMSCs. Our study, therefore, highlights the potential importance of ECM HSPGs  
11 as regulators of BMP-2 activity, giving new insights into the molecular basis of ECM-BMP  
12 interactions and opening avenues for novel strategies to design biomimetic materials  
13 functionalized with BMP-2 in regenerative medicine.  
14  
15  
16  
17  
18  
19  
20  
21  
22  
23  
24  
25  
26  
27  
28  
29  
30

## 31 **Experimental Section**

32 Further information on the experimental materials and methods are available in the supporting  
33 information.  
34  
35

## 36 **Supporting Information**

37 Supporting Information is available from the Wiley Online Library.  
38  
39  
40  
41  
42

## 43 **Acknowledgements**

44 We thank Prof. Joachim Spatz (Dept. Biophysical Chemistry, Heidelberg University and Max  
45 Planck Institute for Medical Research, Heidelberg) for his kind support and fruitful  
46 discussions, Dr. Hugues Lortat-Jacob (IBS, Grenoble, France) for providing HS and valuable  
47 feedback on the experiments; Prof. Walter Sebald, (Würzburg University, Germany) for  
48 providing EHBMP-2. Aïseta Baradji (University of Liverpool, UK and CICbiomaGUNE, San  
49 Sebastian, Spain) for helping with the SUV preparation; Dr. Burcu Minsky (Max Planck  
50 Institute for Intelligent Systems, Stuttgart) for support on SE measurements and, *of course*,  
51 the Cell Adhesion and Signalling group (Heidelberg University). This project has received  
52 funding from: European Union's Framework Program for Research and Innovation Horizon  
53 2020 (2014-2020) under the Marie Skłodowska-Curie Grant Agreement No. H2020-MSCA-  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65

1 IF-2014 and from CellNetworks Cluster, Heidelberg University, under the CellNetworks  
2 Postdoctoral Program 2014-2015. EACA greatly acknowledges the support of DFG (SFB  
3 TRR 79 projects M9 and 5). EM, SW, JPS and EACA thank the Max Planck Society for  
4 support. RPR acknowledges the Spanish Ministry for economy and competitiveness (project  
5 MAT2014-54867-R).  
6

7 Received: ((will be filled in by the editorial staff))

8 Revised: ((will be filled in by the editorial staff))

9 Published online: ((will be filled in by the editorial staff))  
10

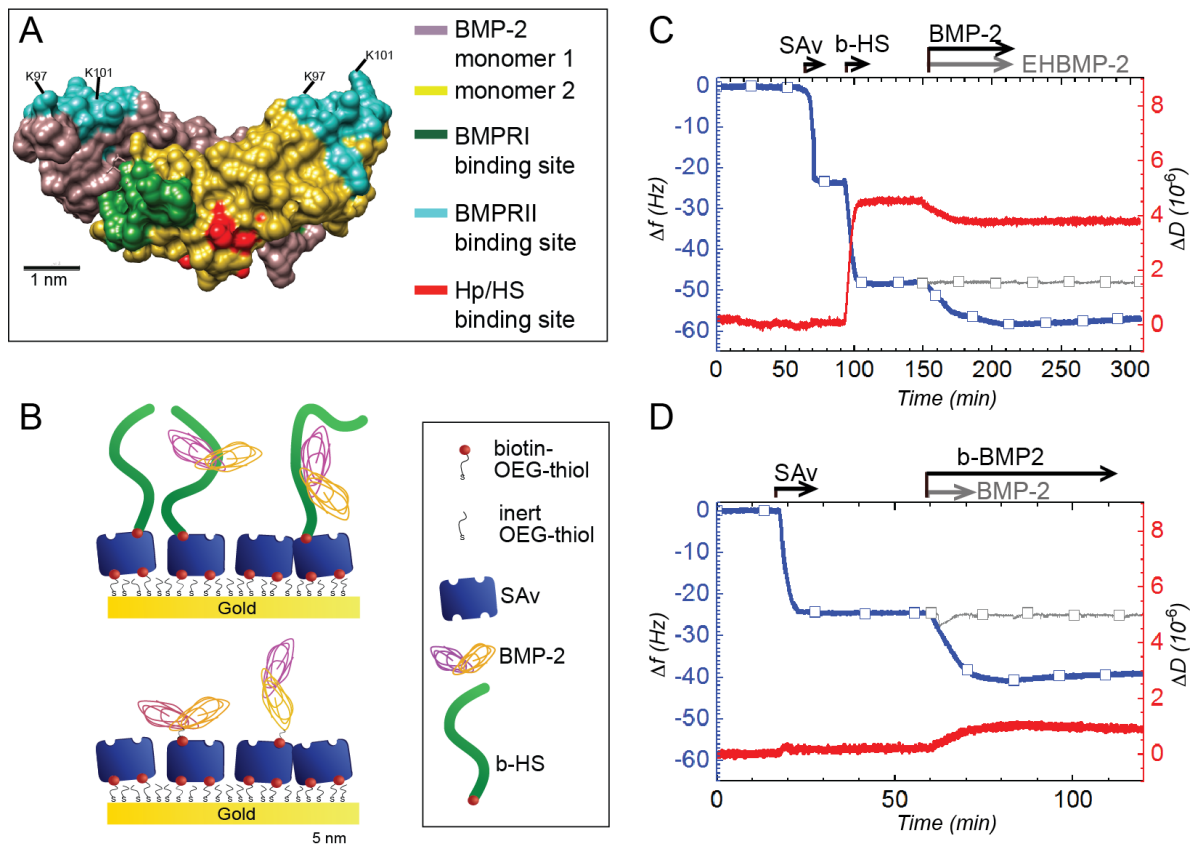
### 11 Reference:

- 12
- 13 [1] H. M. Ryoo, M. H. Lee, Y. J. Kim, *Gene*. **2006**. 366, 51.
- 14
- 15 [2] T. Katagiri, A. Yamaguchi, M. Komaki, E. Abe, N. Takahashi, T. Ikeda, V. Rosen, J.  
16 M. Wozney, A. Fujisawa-Sehara, T. Suda, *J Cell Biol*. **1994**. 127, 1755.
- 17
- 18 [3] A. Asakura, M. Komaki, M. Rudnicki, *Differentiation*. **2001**. 68, 245.
- 19
- 20 [4] F. Liu, A. Hata, J. C. Baker, J. Doody, J. Carcamo, R. M. Harland, J. Massague,  
21 *Nature*. **1996**. 381, 620.
- 22
- 23 [5] M. Geiger, R. H. Li, W. Friess, *Adv Drug Deliv Rev*. **2003**. 55, 1613.
- 24
- 25 [6] B. Sedaghati, B. Hoyer, A. Aigner, M.C. Hacker, M. Schulz-Siegmund, *Regenerative*  
26 *Medicine - from Protocol to Patient: 3. Tissue Engineering, Biomaterial and Nanotechnology*,  
27 Vol. 3, Springer, **2016**.
- 28
- 29 [7] S. Ahmed, R. P. Metpally, S. Sangadala, B. V. Reddy, *J Mol Graph Model*. **2010**. 28,  
30 670.
- 31
- 32 [8] E. Migliorini, A. Valat, C. Picart, E. A. Cavalcanti-Adam, *Cytokine Growth Factor*  
33 *Rev*. **2016**. 27, 43.
- 34
- 35 [9] M. M. Martino, J. A. Hubbell, *Faseb j*. **2010**. 24, 4711.
- 36
- 37 [10] X. Jiao, P. C. Billings, M. P. O'Connell, F. S. Kaplan, E. M. Shore, D. L. Glaser, *J*  
38 *Biol Chem*. **2007**. 282, 1080.
- 39
- 40 [11] D. S. Bramono, S. Murali, B. Rai, L. Ling, W. T. Poh, Z. X. Lim, G. S. Stein, V.  
41 Nurcombe, A. J. van Wijnen, S. M. Cool, *Bone*. **2012**. 50, 954.
- 42
- 43
- 44
- 45
- 46
- 47
- 48
- 49
- 50
- 51
- 52
- 53
- 54
- 55
- 56
- 57
- 58
- 59
- 60
- 61
- 62
- 63
- 64
- 65

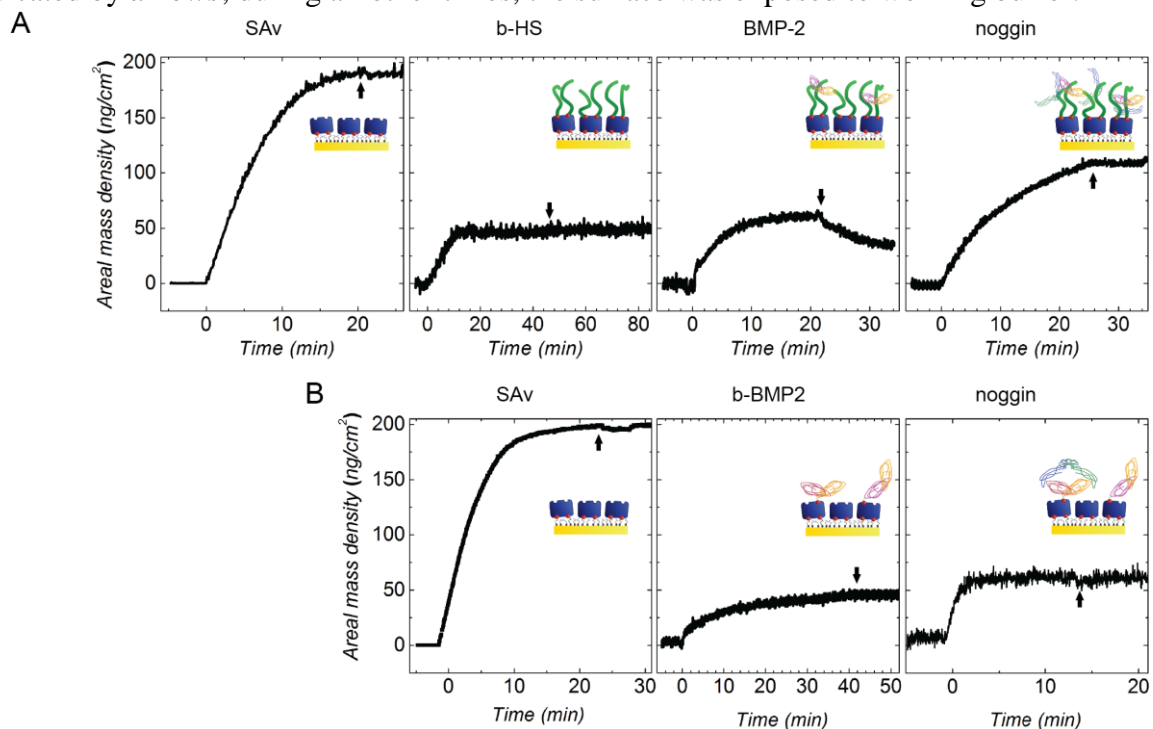
- 1 [12] R. Ruppert, E. Hoffmann, W. Sebald, *Eur J Biochem.* **1996.** 237, 295.
- 2 [13] I. Matsuo, C. Kimura-Yoshida, *Philos Trans R Soc Lond B Biol Sci.* **2014.** 369.
- 3 [14] M. C. Fisher, Y. Li, M. R. Seghatoleslami, C. N. Dealy, R. A. Kosher, *Matrix Biol.*
- 4 **2006.** 25, 27; J. Huegel, M. Enomoto-Iwamoto, F. Sgariglia, E. Koyama, M. Pacifici, *Am J*
- 5 *Pathol.* **2015.** 185, 1676.
- 6 [15] W. J. Kuo, M. A. Digman, A. D. Lander, *Mol Biol Cell.* **2010.** 21, 4028.
- 7 [16] B. L. Viviano, S. Paine-Saunders, N. Gasiunas, J. Gallagher, S. Saunders, *J Biol Chem.*
- 8 **2004.** 279, 5604.
- 9 [17] J. Groppe, J. Greenwald, E. Wiater, J. Rodriguez-Leon, A. N. Economides, W.
- 10 Kwiatkowski, M. Affolter, W. W. Vale, J. C. Izpisua Belmonte, S. Choe, *Nature.* **2002.** 420,
- 11 636.
- 12 [18] G. Sanchez-Duffhues, C. Hiepen, P. Knaus, P. Ten Dijke, *Bone.* **2015.** 80, 43; Y.
- 13 Matsumoto, K. Matsumoto, F. Irie, J. Fukushi, W. B. Stallcup, Y. Yamaguchi, *J Biol Chem.*
- 14 **2010.** 285, 19227.
- 15 [19] R. Gomes, C. Kirn-Safran, M. C. Farach-Carson, D. D. Carson, *J Musculoskelet*
- 16 *Neuronal Interact.* **2002.** 2, 511.
- 17 [20] S. Murali, B. Rai, C. Dombrowski, J. L. Lee, Z. X. Lim, D. S. Bramono, L. Ling, T.
- 18 Bell, S. Hinkley, S. S. Nathan, J. H. Hui, H. K. Wong, V. Nurcombe, S. M. Cool,
- 19 *Biomaterials.* **2013.** 34, 5594.
- 20 [21] S. Sarrazin, W. C. Lamanna, J. D. Esko, *Cold Spring Harb Perspect Biol.* **2011.** 3.
- 21 [22] D. Thakar, E. Migliorini, L. Coche-Guerente, R. Sadir, H. Lortat-Jacob, D. Boturyn, O.
- 22 Renaudet, P. Labbe, R. P. Richter, *Chem Commun.* **2014.** 50, 15148.
- 23 [23] E. Migliorini, D. Thakar, R. Sadir, T. Pleiner, F. Baleux, H. Lortat-Jacob, L. Coche-
- 24 Guerente, R. P. Richter, *Biomaterials.* **2014.** 35, 8903.
- 25 [24] W. A. Hendrickson, A. Pahler, J. L. Smith, Y. Satow, E. A. Merritt, R. P. Phizackerley,
- 26 *Proc Natl Acad Sci U S A.* **1989.** 86, 2190.

- 1  
2  
3  
4  
5  
6  
7  
8  
9  
10  
11  
12  
13  
14  
15  
16  
17  
18  
19  
20  
21  
22  
23  
24  
25  
26  
27  
28  
29  
30  
31  
32  
33  
34  
35  
36  
37  
38  
39  
40  
41  
42  
43  
44  
45  
46  
47  
48  
49  
50  
51  
52  
53  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65
- [25] N. S. Gandhi, R. L. Mancera, *Biochim Biophys Acta*. **2012**. 1824, 1374.
- [26] E. Migliorini, D. Thakar, J. Kuhnle, R. Sadir, D. P. Dyer, Y. Li, C. Sun, B. F. Volkman, T. M. Handel, L. Coche-Guerente, D. G. Fernig, H. Lortat-Jacob, R. P. Richter, *Open Biol*. **2015**. 5.
- [27] J. T. Gallagher, J. E. Turnbull, M. Lyon, *Int J Biochem*. **1992**. 24, 553.
- [28] C. I. Gama, S. E. Tully, N. Sotogaku, P. M. Clark, M. Rawat, N. Vaidehi, W. A. Goddard, 3rd, A. Nishi, L. C. Hsieh-Wilson, *Nat Chem Biol*. **2006**. 2, 467.
- [29] N. Jastrebova, M. Vanwildemeersch, U. Lindahl, D. Spillmann, *J Biol Chem*. **2010**. 285, 26842; M. Maccarana, Y. Sakura, A. Tawada, K. Yoshida, U. Lindahl, *J Biol Chem*. **1996**. 271, 17804.
- [30] L. Luca, M. A. Capelle, G. Machaidze, T. Arvinte, O. Jordan, R. Gurny, *Int J Pharm*. **2010**. 391, 48.
- [31] S. Masuda, K. Namba, H. Mutai, S. Usui, Y. Miyanaga, H. Kaneko, T. Matsunaga, *Biochem Biophys Res Commun*. **2014**. 447, 496.
- [32] D. Yaffe, O. Saxel, *Nature*. **1977**. 270, 725.
- [33] T. Katagiri, S. Akiyama, M. Namiki, M. Komaki, A. Yamaguchi, V. Rosen, J. M. Wozney, A. Fujisawa-Sehara, T. Suda, *Exp Cell Res*. **1997**. 230, 342.
- [34] E. H. Schwab, T. L. Pohl, T. Haraszti, G. K. Schwaerzer, C. Hiepen, J. P. Spatz, P. Knaus, E. A. Cavalcanti-Adam, *Nano Lett*. **2015**. 15, 1526.
- [35] T. L. Pohl, J. H. Boergemann, G. K. Schwaerzer, P. Knaus, E. A. Cavalcanti-Adam, *Acta Biomater*. **2012**. 8, 772; P. Paarmann, G. Dorpholz, J. Fiebig, A. R. Amsalem, M. Ehrlich, Y. I. Henis, T. Muller, P. Knaus, *Int J Biochem Cell Biol*. **2016**. 76, 51; T. Takada, T. Katagiri, M. Ifuku, N. Morimura, M. Kobayashi, K. Hasegawa, A. Ogamo, R. Kamijo, *J Biol Chem*. **2003**. 278, 43229.
- [36] B. Mulloy, C. Gee, S. F. Wheeler, R. Wait, E. Gray, T. W. Barrowcliffe, *Thromb Haemost*. **1997**. 77, 668.

- 1 [37] R. Richter, A. Mukhopadhyay, A. Brisson, *Biophys J.* **2003.** 85, 3035.  
2  
3 [38] N. B. Eisele, S. Frey, J. Piehler, D. Gorlich, R. P. Richter, *EMBO Rep.* **2010.** 11, 366.  
4  
5 [39] I. Carton, A. R. Brisson, R. P. Richter, *Anal Chem.* **2010.** 82, 9275.  
6  
7 [40] G. V. Dubacheva, T. Curk, B. M. Mognetti, R. Auzely-Velty, D. Frenkel, R. P. Richter,  
8  
9 *J Am Chem Soc.* **2014.** 136, 1722.  
10  
11 [41] M. B. Huglin, *Journal of Applied Polymer Science.* **1965.** 9, 1097.  
12  
13 [42] R. P. Richter, K. B. Rodenhausen, E. N. B., M. Schubert, *Coupling Spectroscopic*  
14  
15 *Ellipsometry and Quartz Crystal Microbalance to Study Organic Films at the Solid-Liquid*  
16  
17 *Interface*, Vol. 52, Springer, Heidelberg **2014.**  
18  
19 [43] V. Hagel, T. Haraszti, H. Boehm, *Biointerphases.* **2013.** 8, 36; T. T. Tsay, K. A.  
20  
21 Jacobson, *Biophys J.* **1991.** 60, 360.  
22  
23 [44] W. Wagner, P. Horn, M. Castoldi, A. Diehlmann, S. Bork, R. Saffrich, V. Benes, J.  
24  
25 Blake, S. Pfister, V. Eckstein, A. D. Ho, *PLoS One.* **2008.** 3, e2213.  
26  
27 [45] J. Kopf, P. Paarmann, C. Hiepen, D. Horbelt, P. Knaus, *Biofactors.* **2014.** 40, 171.  
28  
29  
30  
31  
32  
33  
34  
35  
36  
37  
38  
39  
40  
41  
42  
43  
44  
45  
46  
47  
48  
49  
50  
51  
52  
53  
54  
55  
56  
57  
58  
59  
60  
61  
62  
63  
64  
65

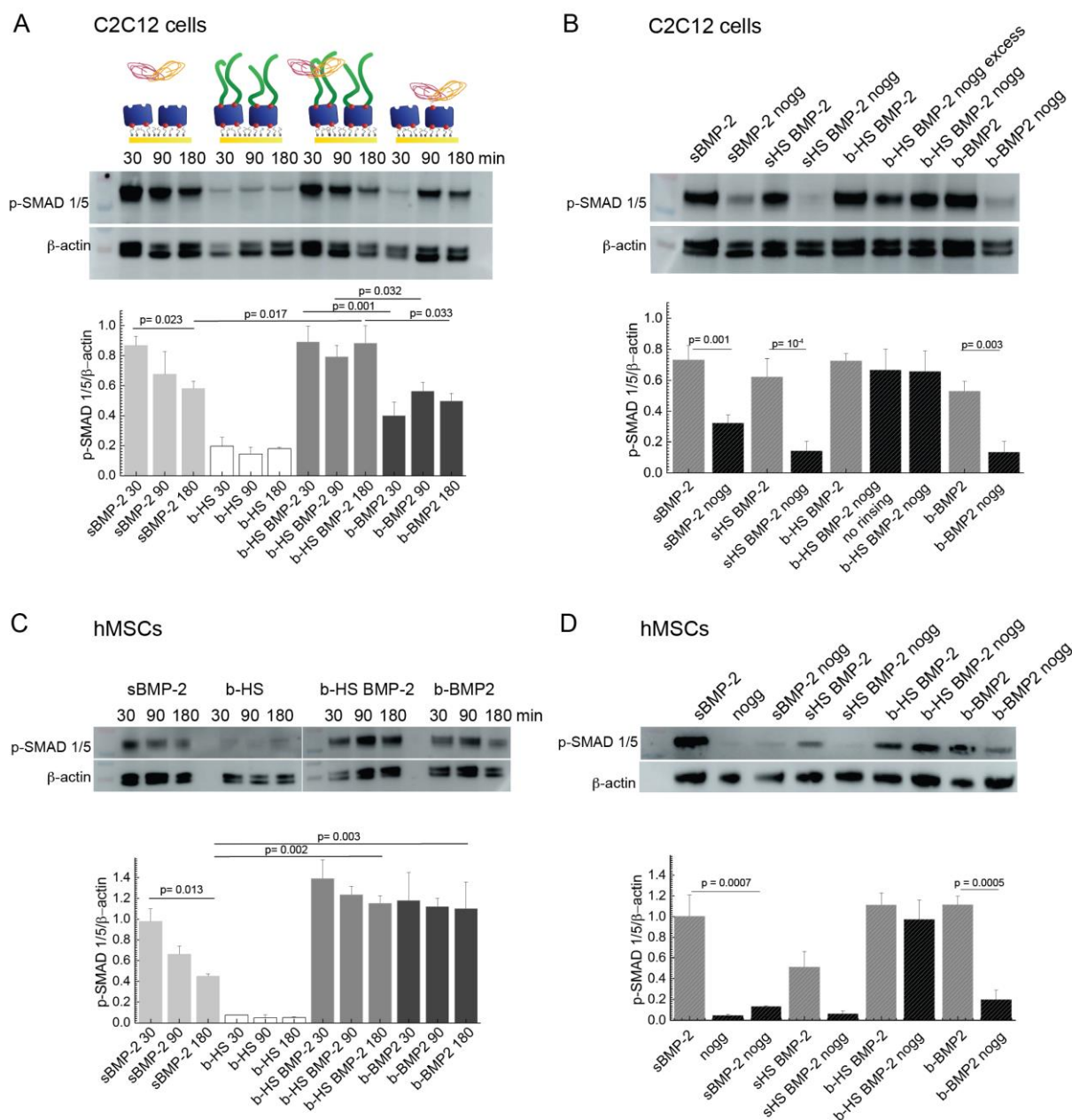


**Figure 1.** **A:** BMP-2 homodimer structure (PDB: 3BMP). **B:** Schematic representation of the biomimetic platforms. **C-D:** QCM-D characterization. Frequency shifts:  $\Delta f$  - blue lines with square symbols; dissipation shifts:  $\Delta D$  - red lines. Start and duration of incubation steps are indicated by arrows; during all other times, the surface was exposed to working buffer.





**Figure 2.** Surface functionalization is followed *in situ* by SE on a gold-supported OEG monolayer. Sample concentrations are: 83 nM SAV, 0.8  $\mu$ M b-HS, 96 nM BMP-2, b-BMP2 and 200 nM noggin. Each incubation step started at 0 min; start of rinsing in working buffer is indicated by arrows.



**Figure 3.** C2C12 cells (**A, B**) and hMSCs (**C, D**) plated on functionalized surfaces and lysed, at different time points (**A, C**) or after 90 minutes on substrates incubated with or without noggin (**B, D**). p-SMAD 1/5 expression analyzed by Western Blot.

**Table 1.** Data was extracted from SE measurements at signal stabilization (**Figure 2**). Mean values and standard errors from the mean were derived from 3 independent measurements. On the basis of these numbers we quantify the number of molecules grafted per unit surface area and the stoichiometry of binding.

Compound	Areal mass density (ng/cm <sup>2</sup> )
SAv	204.3±9.9
+ b-BMP2	44.2±4.6
+ noggin	57.6±2.1
b-HS	40.7±3.5
+ BMP-2 <sub>(eq)</sub>	53.2±4.6
+ BMP-2 <sub>(rins)</sub>	35.3±2.7
+ Noggin	112±4.1

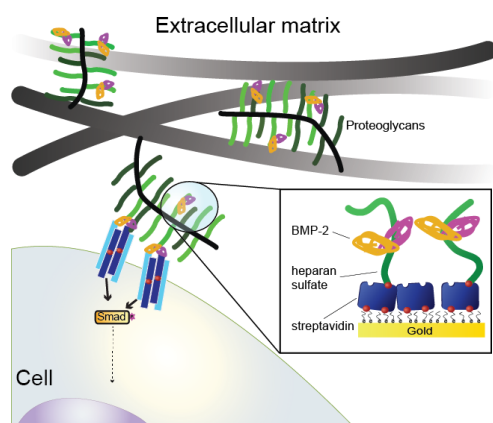
<sup>(eq)</sup> value close to equilibrium during BMP-2 injection; <sup>(rins)</sup> value after BMP-2 rinsing with working buffer until plateau was reached.

**Table of contents: Biomimetic platforms reproducing *in vitro* the BMP-2 presentation via extracellular matrix (ECM)-associated heparan sulfate proteoglycans (HSPGs).** A bottom-up approach is used to functionalize biomimetic surfaces and to investigate the role of ECM-associated HS on the biological activity of BMP-2.

**Keyword:** BMP-2, heparan sulfate, extracellular matrix, noggin, biomimetic platforms

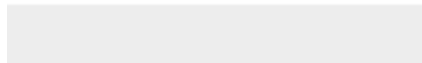
*Elisa Migliorini\*, Patrick Horn, Tamás Haraszti, Seraphine V. Wegner, Christian Hiepen, Petra Knaus, Ralf P. Richter and E. Ada Cavalcanti-Adam\**

**Title: Enhanced biological activity of BMP-2 bound to surface-grafted heparan sulfate**





Click here to access/download  
**Supporting Information**  
Supporting Information.pdf





Click here to access/download  
**Supporting Information**  
FRAPAnalysis