

Supporting information

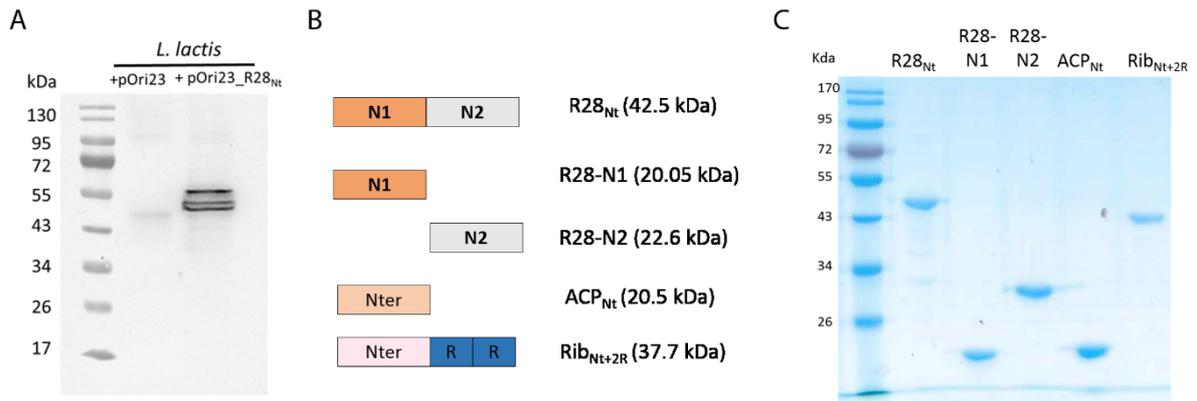


Figure S1. Schematic representation of constructed peptides used in this study, and their expression in *L. lactis* or *E. coli*. *A*, Western blot with anti-R28_{Nt} antiserum of cell wall extracts of *L. lactis* expressing R28_{Nt}, + pOri_R28_{Nt}; or not, +pOri. *B*, schematic representation of the different peptides used in the study. Color code corresponds to Fig. 1A. *C*, Coomassie blue staining of an acrylamide gel with 1 μg of each peptide loaded.

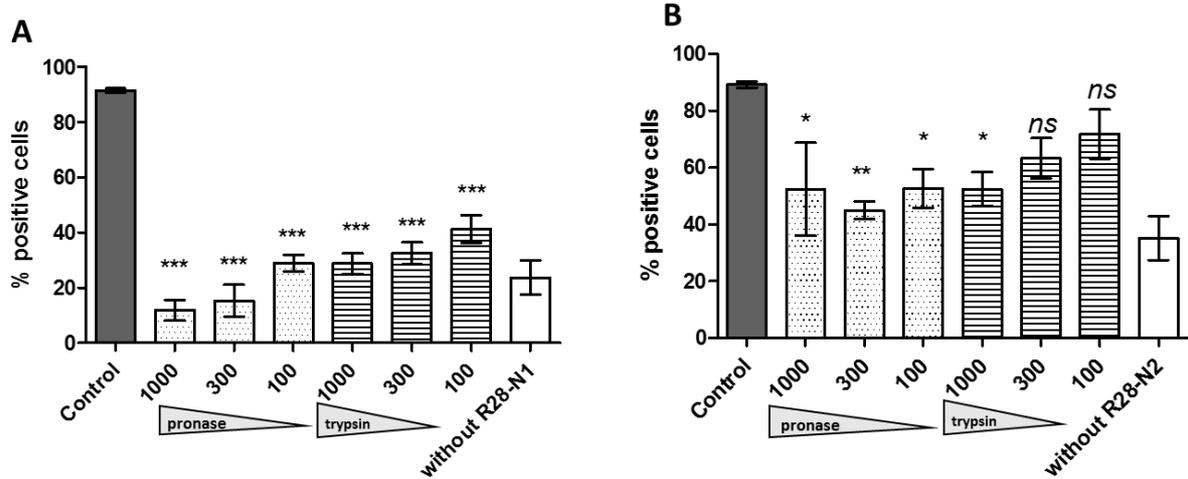


Figure S2. Effect of different treatments on the binding of A, R28-N1 and B, R28-N2 to HEC-1-A cells. Treatments were applied to HEC-1-A cells in suspension prior to incubation with purified R28-N1 and R28-N2; bound peptides were immunolabelled and cells were analyzed by flow cytometry. Statistical analysis was performed against the untreated condition Error bars correspond to SEM of three independent experiments. Two-Way ANOVA: *, $p < 0.05$; **, $p < 0.005$; ***, $p < 0.001$; ns, not significant.

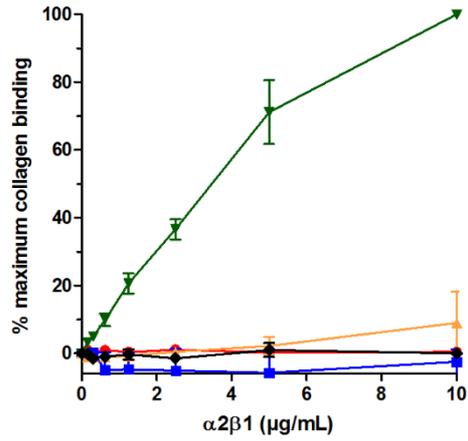


Figure S3. R28Nt, R28-N1 and R28-N2 do not bind the $\alpha 2\beta 1$ integrin. Assessment of integrin $\alpha 2\beta 1$ binding to R28_{Nt}, red; R28-N1, blue; R28-N2, orange; Type I collagen, green; BSA, black. The binding is expressed as percentage of that of the collagen at 10 $\mu\text{g/mL}$ integrin.

Table S1. Strains and plasmids used in this study

Strains or plasmid	Relevant properties	Source or reference
<u><i>Escherichia coli</i></u>		
BL21 λDE3	F ⁻ <i>ompT gal (dcm) (lon) hsdSB (rB⁻ mB⁻) endA1 hsdR17(rK⁻mK⁺)</i>	(1)
<u><i>Streptococcus pyogenes</i></u>		
M28PF1	<i>emm28</i> source of DNA for R28 _{Nt} , R28-N1, R28-N2 amplifications	(2)
ΔR28	M28PF1 lacking the <i>Spy1336</i> coding for R28	This study
<u><i>Streptococcus agalactiae</i></u>		
A909	serotype Ia, sequence type 7 , used for amplification of the gene of ACP	(3)
BM110	serotype III, sequence type 17, used for the amplification of the gene encoding Rib	(4)
<u><i>Lactococcus lactis</i></u>		
MG1363	Lac ⁻ Prt ⁻ ; NCDO 712 derivative	(5)
CCH2022	MG1363+ pOri23	This study
CCH2023	MG1363+ pOri23_R28 _{Nt}	This study
<u>Plasmids</u>		
pG+host5	Erm; ColE1 replicon, thermosensitive derivative of pGK12; MCS pBluescript	(6)
pG1_R28	500 pb upstream (R28F1-R28R1) and 500 bp downstream (R28F2-R28R2) sequence of <i>Spy1336</i> cloned in pG+host5	This study
pET2818	Amp, oriR pBR322, T7 promoter, His-Tag coding sequence, pET28/16 derivative.	S. Mesnage (Pers. com.)
pET2818_R28 _{Nt}	pET2818 with the sequence encoding the R28 _{Nt} , without signal peptide. For the production of the peptide in <i>E. coli</i> with a His Tag	This study
pET2818_R28.N1	Ibidem with the sequence encoding R28-N1	This study
pET2818_R28.N2	Ibidem with the sequence encoding R28-N2	This study
pET2818_ACP	Ibidem with the sequence encoding ACP	This study
pOri23	Erm; <i>ermB</i> , ori ColE1, thermosensitive derivative of pIL253, P23 promoter of <i>L. lactis</i> MG1363	(7)
pOri23_R28 _{Nt}	High level of expression of R28 _{Nt} at the cell wall of Gram-positive bacteria	This study

Table S2. Primers used in this study for cloning and protein purifications

Primer Name	Sequence*
R28-F1	CGACTCTAGAG GATCC GCCTGTGAGAGACGATCATA
R28-R1	ATCATTCTTATCGGCCCGTTGTTTCGTCTGTGAAG
R28-F2	GCCGATAAGAATGATCCAGC
R28-R2	CCATGATTAC GAA TT CGC CTTAACTCGTATTCCTTT
R28_F	CCATGG CTACAATTCCAGGGAGTGC
R28_R	GGATCC ACCTTTTGGTTCGTTCGTATC
R28_N1_F	CCATGG GGTCTACAATTCC
R28_N1_R	GGATC CTGGTAGCGATAACAATAA
R28_N2_F	CCATGG ATAAAATTAAGTATTCGCC
R28_N2_R	GGATC CTTTTGGTTCGTTG
ACP_F	CCATGG GGTCAATAGTTGCTGCATCTACAAT
ACP_R	GGATC CTACTGACAATACTAACAATTTCTC

* restriction enzyme sites used for cloning are highlighted in bold

Supporting information references:

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6. Biswas, I., Gruss, A., Ehrlich, S. D., and Maguin, E. (1993) High-efficiency gene inactivation and replacement system for gram-positive bacteria. *J. Bacteriol.* **175**, 3628–35
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Supporting information file

The raw data of hits found by mass-spectrometry analysis of the co-immunoprecipitations are available in the supporting information file. In green and yellow are highlighted hits that are significantly enriched in the presence of R28_{Nt} in all three independent experiments. In yellow are the surface exposed membrane proteins that correspond to Table 1 hits. In pink are highlighted the samples in which the given protein was not detected. CT-1, CT-2, CT-3 and R28-1, R28-2, R28-3, samples eluted from the control zones and R28_{Nt} co-immunoprecipitated zones of experiments 1, 2, 3, respectively