

**Genipin crosslinking reduced the immunogenicity of xenogeneic decellularized porcine whole-liver matrices through regulation of immune cell proliferation and polarization**

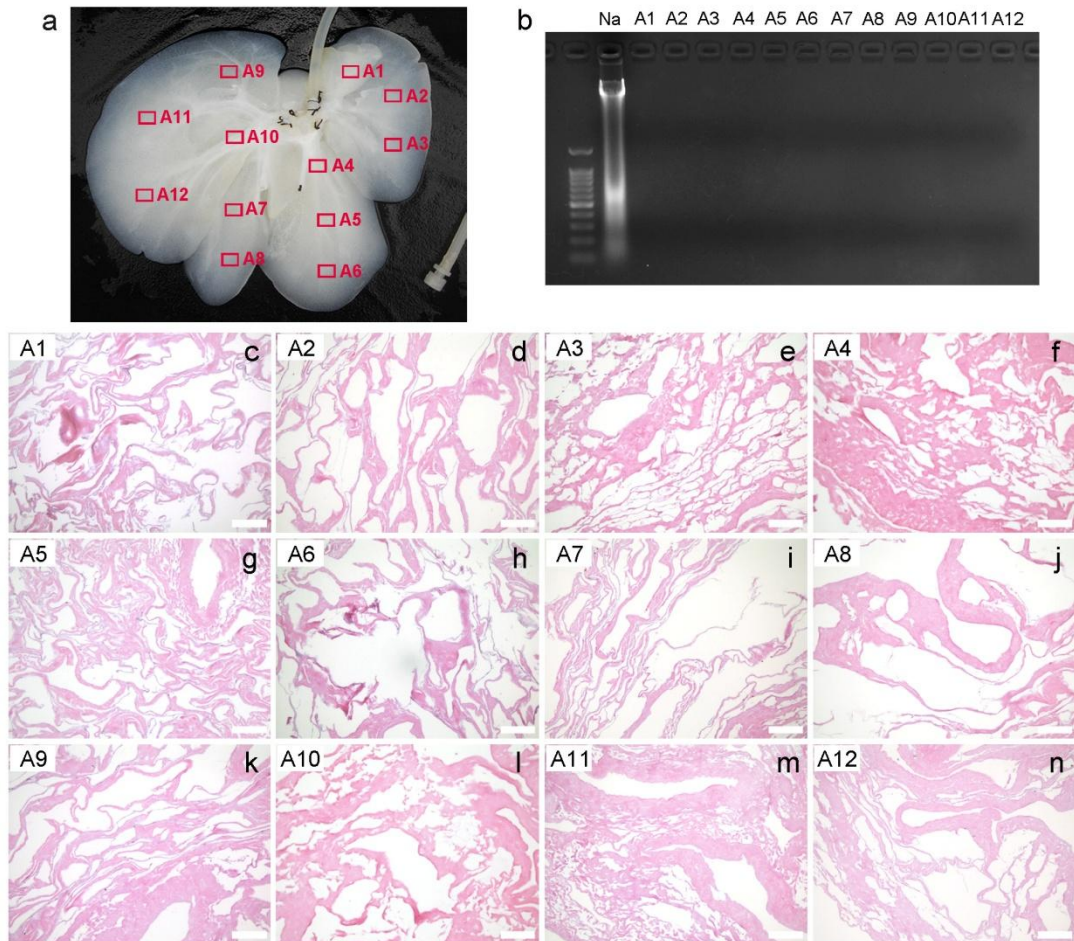
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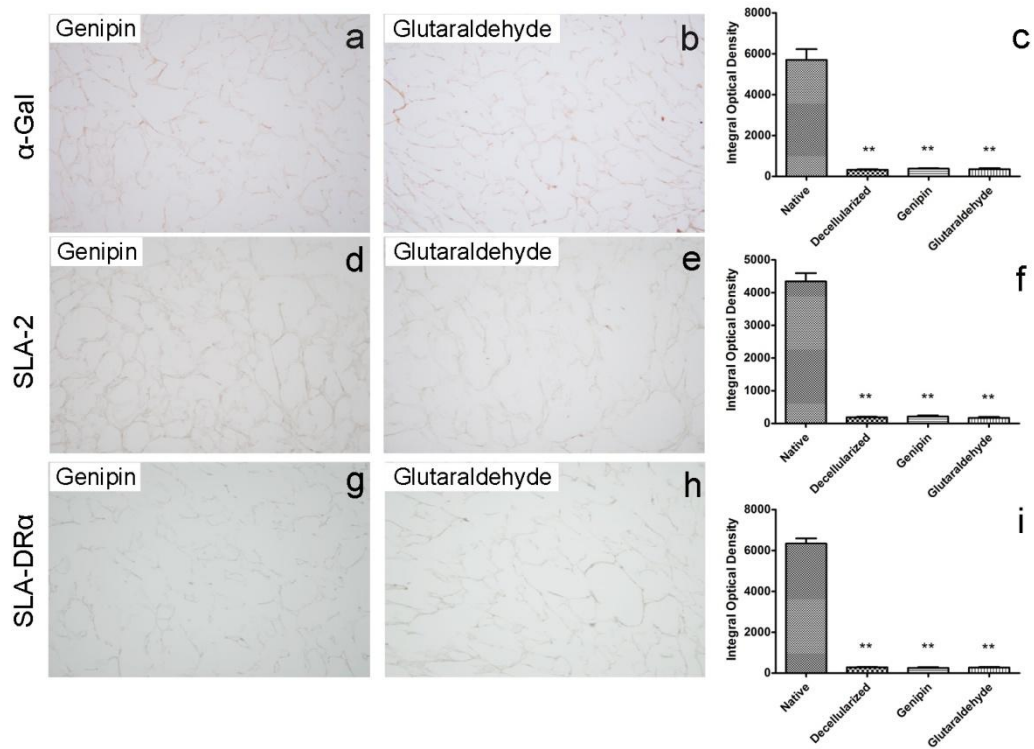
## Supplementary Information



**Supplementary Fig. S1** Analysis of decellularization uniformity.

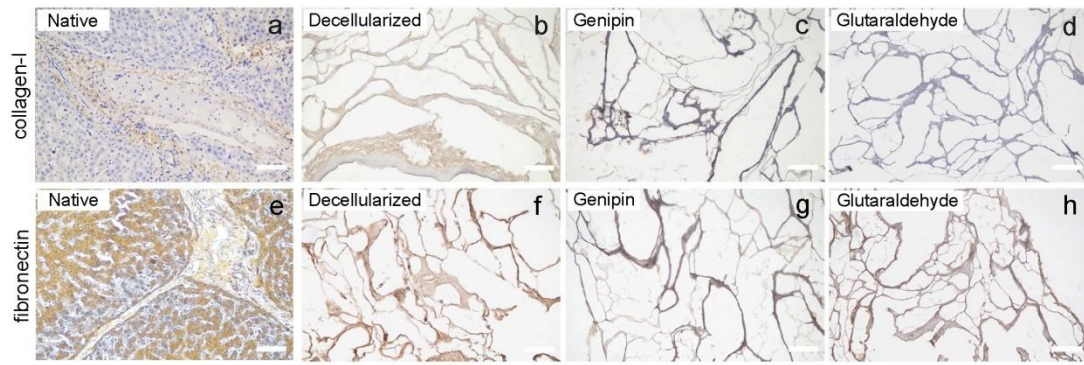
(a) The morphology of decellularized liver and the indicated 12 different areas. (b) Agarose gel electrophoresis of DNA extracted from selected decellularized liver matrices. Na= native control. (c-n) Histological images of selected liver ECM stained with H&E indicating decellularization uniformity.

Scale bars = 100  $\mu$ m.



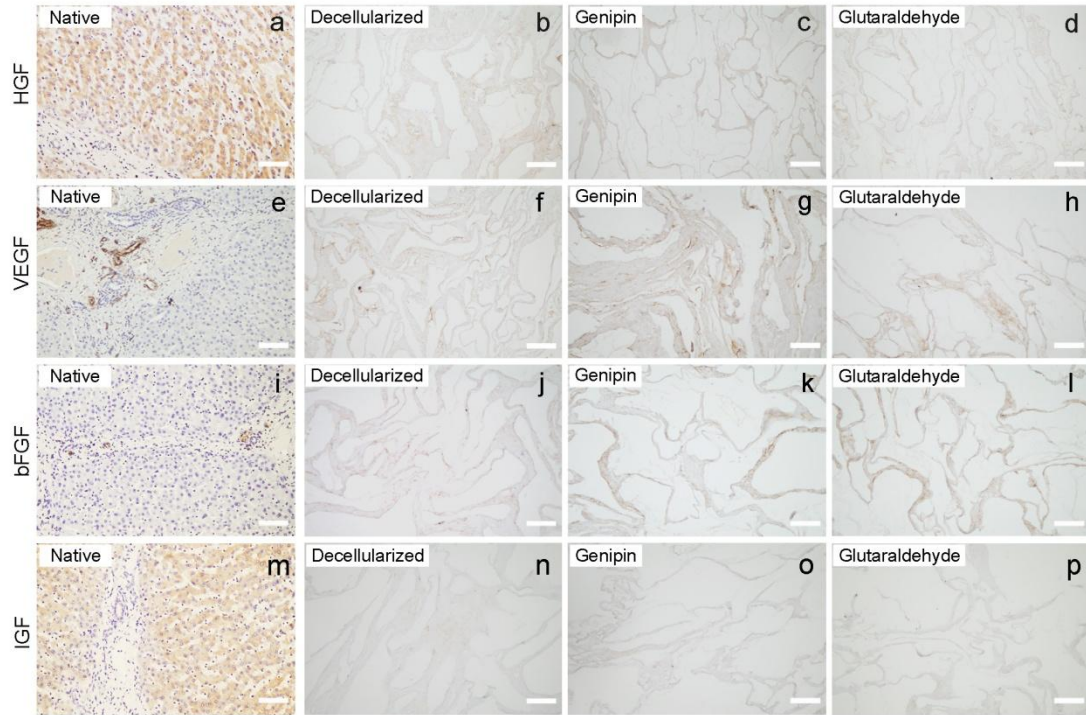
**Supplementary Fig. S2** Immunohistochemistry and quantification of immunogenic or pathogenic antigens on matrix.

Immunohistochemical staining of genipin and glutaraldehyde crosslinked liver ECMs for  $\alpha$ -Gal (a, b), SLA-2 (d, e), SLA-DR $\alpha$  (g, h) are shown. Scale bars = 100  $\mu$ m. Immunohistochemical quantification of  $\alpha$ -Gal (c), SLA-2 (f), SLA-DR $\alpha$  (i) are also shown. \*\* $p < 0.01$  with respect to the native group. All data are given as the mean  $\pm$  SEM.



**Supplementary Fig. S3** Immunohistochemistry of matrix components.

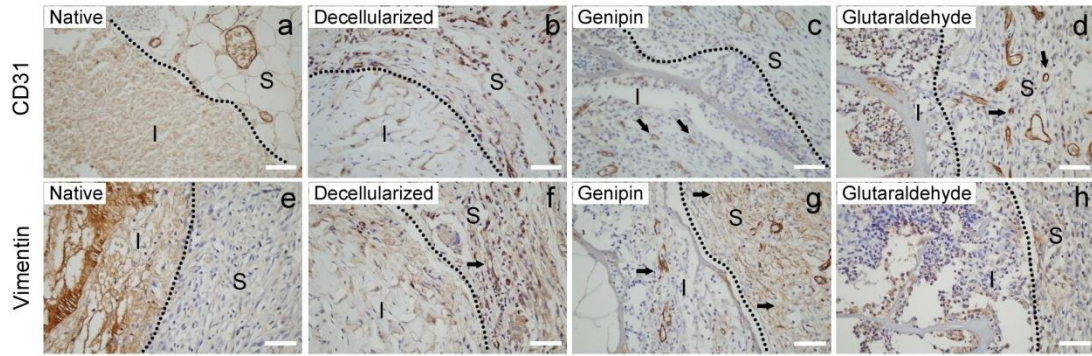
(a–p) Immunohistochemistry staining of native liver and un-crosslinked and crosslinked decellularized liver ECM for collagen I and fibronectin. Scale bars = 100  $\mu\text{m}$ .



**Supplementary Fig. S4** Immunohistochemistry of residual growth factors in liver matrix.

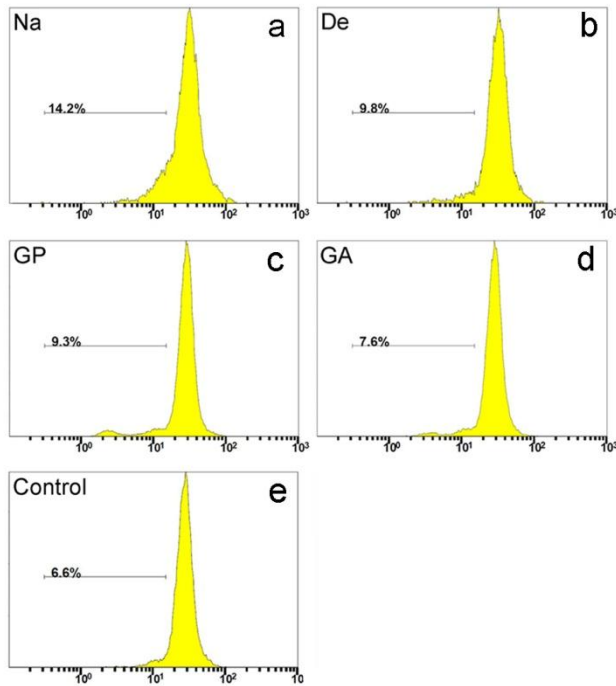
(a–p) Immunohistochemistry staining of native liver, un-crosslinked and crosslinked liver ECM for

HGF, VEGF, bFGF, and IGF (from top to bottom). Scale bars = 100 μm.



**Supplementary Fig. S5** Immunohistochemistry of the liver xenografts.

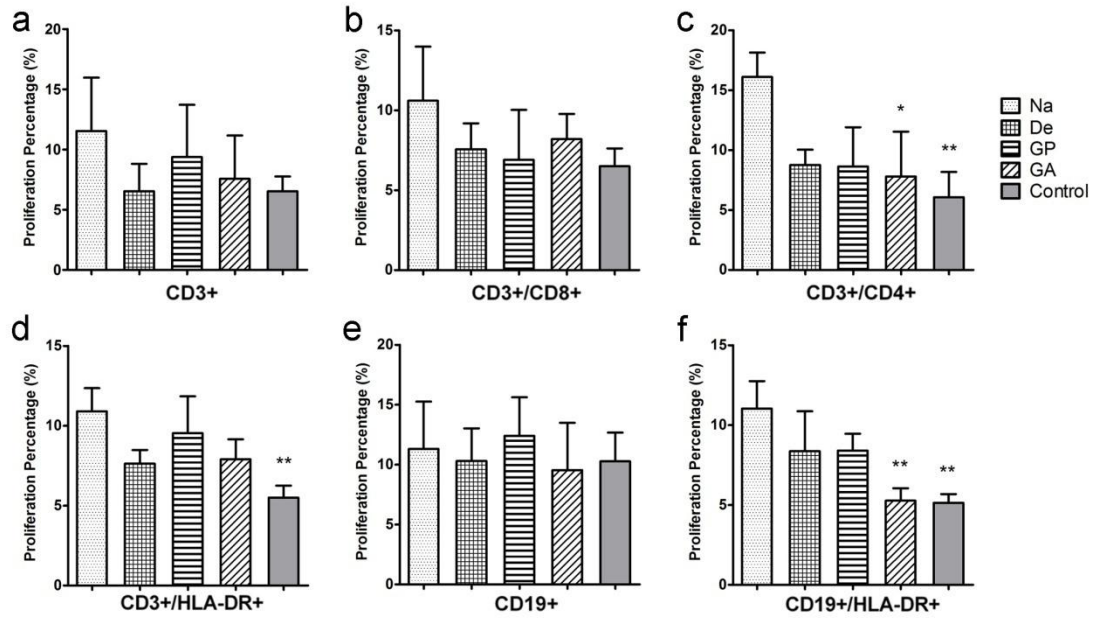
(a-h) Immunohistochemistry towards CD31 and vimentin were examined in native liver, decellularized liver ECM, genipin-fixed ECM, and glutaraldehyde-fixed ECM at 21 days post-surgery to show the vascular endothelial cells (arrowed) and fibroblast cells (arrowed). Scale bars = 50  $\mu\text{m}$ . The dotted line indicates the border of the implants and surrounding tissue. Abbreviations: S=surrounding tissue; I=implanted porcine liver materials.



**Supplementary Fig. S6** Proliferation properties of PBMCs in co-culture with liver matrices without OKT3.

(a-e) Representative FACS histograms of immune cells co-cultured without OKT3 stimulus.

Proliferation responses without protein extracts were used as a negative control.



**Supplementary Fig. S7** Impact of porcine liver matrices on T cell and B cell subpopulation

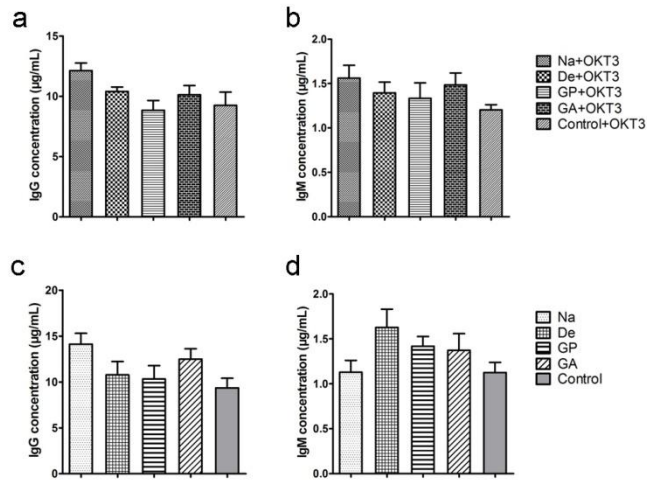
proliferation in co-cultures without CD3 stimulus.

(a-f) The proliferation patterns of T cells, B cells and their subsets was analyzed using anti-human CD3,

CD8, CD4, HLA-DR, CD19 antibodies. \* $p < 0.05$  with respect to Na group, \*\* $p < 0.01$  with respect to

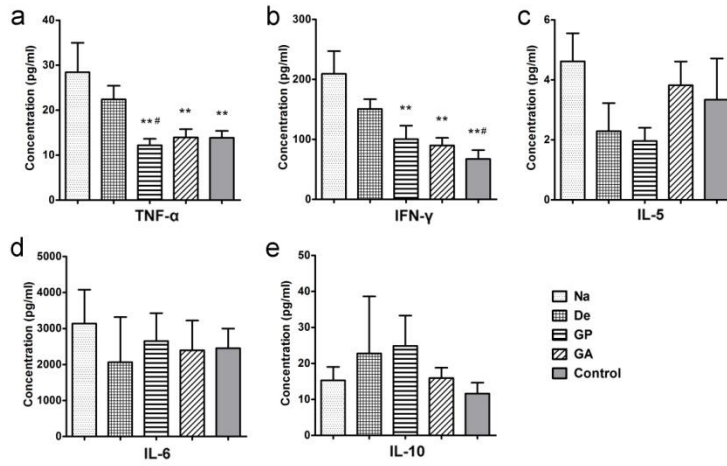
Na group. All data are given as the mean  $\pm$  SEM (n=3).





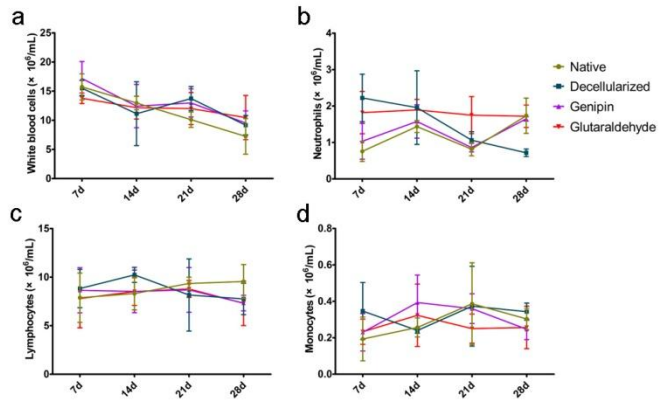
**Supplementary Fig. S8** Impact of porcine liver matrices on B cell activation *in vitro*.

IgG and IgM production in human PBMC co-culture with liver material extracts in the presence of OKT3 (a,b) or without OKT3 (c,d).



**Supplementary Fig. S9** Th1/Th2 cytokine secretion profile of PBMCs co-cultured with protein extracts and without anti-CD3 stimulus.

(a-e) Th1/Th2 cytokine levels are shown for co-cultures of PBMCs alone or in combination with protein extracts of porcine matrix. \*\*p<0.01 with respect to native group, #p<0.05 with respect to De group. All data are given as the mean ± SEM (n=3).



**Supplementary Figure. S10** Systemic white blood cell counts of host rats.

(a-d) total white blood cell count, neutrophil count, lymphocyte count, and monocyte count are shown,

indicating no differences among groups. All data are given as the mean  $\pm$  SEM (n=3).