Supporting Information

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Fig. S1. Tissue to muscle ratios for iodine-125 (125 I) -labeled peptides p1-p7 in reactive amyloidosis (AA) amyloid-bearing mice (red) or WT littermate control animals (blue) at 1 h postinjection (pi) of radiotracer. Tissue to muscle ratios were calculated using the mean percent injected dose (ID) per gram values (n = 3) for each tissue relative to the mean percent ID per gram values for muscle.



Fig. 52. The ¹²⁵I-p5 peptide binds apolipoprotein A2c amyloid in vivo in the intestines of transgenic mice. Transgenic mice expressing the mouse apolipoprotein A2c allele were injected with ~100 μ Ci ¹²⁵I-p5 peptide and euthanized 2 h pi. Tissues were harvested at necropsy, formalin-fixed tissue sections were prepared, and microautoradiographs were generated as described in *Materials and Methods*. Serial tissue sections were also stained with H&E or Congo red, and images were acquired as described in *Materials and Methods*. (Magnification: 80×.) Although variable in concentration, the presence of amyloid-bound ¹²⁵I-p5 was evidenced in the microautoradiographs as punctate black deposits that were intense in certain villi, and in each case, they colocalized with the red-gold amyloid deposits seen in Congo red-stained tissue section of the intestines.



Fig. S3. Immunohistochemical staining of heparan sulfate. Human amyloid-containing tissues from patients with Alzheimer's disease (*A* and *B*), light chain amyloidosis (*C* and *D*), or AA (*E*) were stained using an antiheparan sulfate monoclonal antibody (F58-10E4 clone; Siekagaku Biobusiness Corp) followed by the addition of a biotinylated secondary reagent; finally, they were detected by the addition of streptavidin-conjugated HRP and visualized using diaminobenzidene (brown coloration). The specific binding of the antiheparan sulfate antibody (arrows) correlated with the distribution of amyloid deposits evidenced in Congo red-stained tissue sections. All slides were counterstained with H&E. (Magnification: 80×.) For display, the images were scaled to 13% of original size.

Table S1. Percent ID per gram ¹²⁵I-peptide in AA mice at 4 h pi (n = 3)

	Peptide percent ID per gram (mean ± SD)								
	p1	p2	р3	p4	р5	p6	р7		
Muscle	0.9 ± 0.3	0.4 ± 0.3	0.5 ± 0.1	0.6 ± 0.1	0.7 ± 0.2	0.5 ± 0.0	0.7 ± 0.3		
Liver	7.6 ± 2.7	1.0 ± 0.5	3.0 ± 0.8	3.2 ± 1.0	5.3 ± 1.0	2.6 ± 0.2	1.9 ± 0.4		
Pancreas	3.9 ± 1.3	0.9 ± 0.7	1.2 ± 0.1	2.1 ± 0.9	9.5 ± 0.9	3.7 ± 1.2	1.8 ± 0.6		
Spleen	7.4 ± 2.8	0.8 ± 0.4	1.0 ± 0.1	3.7 ± 1.1	8.3 ± 1.1	2.8 ± 0.3	1.2 ± 0.2		
L kidney	3.7 ± 0.9	1.7 ± 1.2	1.6 ± 0.4	2.1 ± 0.6	2.7 ± 0.3	1.9 ± 0.0	1.8 ± 0.3		
Stomach	26.2 ± 3.9	7.9 ± 3.3	9.8 ± 2.4	4.6 ± 1.1	8.4 ± 2.4	12.9 ± 2.9	13.0 ± 0.2		
Small Intestine	2.2 ± 0.6	0.8 ± 0.4	1.0 ± 0.3	1.7 ± 0.5	4.0 ± 1.5	1.5 ± 0.6	1.5 ± 0.3		
Large intestine	2.1 ± 0.5	0.6 ± 0.5	1.4 ± 0.2	2.0 ± 0.6	1.9 ± 1.2	1.4 ± 0.2	1.3 ± 0.3		
Heart	2.0 ± 0.4	0.6 ± 0.3	1.1 ± 0.2	1.6 ± 0.3	1.4 ± 0.2	1.0 ± 0.1	1.1 ± 0.2		
Lung	2.8 ± 0.5	1.1 ± 0.5	1.3 ± 0.1	1.6 ± 0.2	1.1 ± 0.0	1.1 ± 0.1	1.5 ± 0.2		
Tongue	nd	nd	1.5 ± 0.1	2.3 ± 0.5	nd	nd	nd		

nd, not determined.

Table S2. Percent ID per gram ¹²⁵I-peptide in WT mice at 4 h pi (n = 3)

	Peptide percent ID/g (mean \pm SD)									
	p1	p2	р3	р4	р5	р6	р7			
Muscle	0.5 ± 0.3	0.1 ± 0.0	0.1 ± 0.1	0.6 ± 0.2	0.2 ± 0.0	0.1 ± 0.1	0.3 ± 0.1			
Liver	1.3 ± 0.3	0.3 ± 0.1	0.3 ± 0.1	2.6 ± 2.3	0.6 ± 0.0	0.5 ± 0.2	1.4 ± 0.2			
Pancreas	0.8 ± 0.5	0.2 ± 0.1	0.2 ± 0.1	1.8 ± 1.0	0.6 ± 0.2	0.6 ± 0.4	0.9 ± 0.4			
Spleen	0.9 ± 0.4	0.2 ± 0.1	0.2 ± 0.0	1.6 ± 0.8	0.4 ± 0.1	0.6 ± 0.2	1.4 ± 0.2			
L kidney	1.2 ± 0.6	0.7 ± 0.2	0.6 ± 0.1	1.5 ± 0.5	1.3 ± 0.1	0.6 ± 0.2	1.6 ± 1.6			
Stomach	9.9 ± 5.3	2.3 ± 0.6	1.4 ± 0.7	6.7 ± 3.3	3.6 ± 1.2	5.2 ± 2.8	6.3 ± 1.0			
Small intestine	1.0 ± 0.4	0.5 ± 0.4	0.3 ± 0.2	1.3 ± 0.6	0.7 ± 0.3	0.5 ± 0.2	0.7 ± 0.2			
Large intestine	0.7 ± 0.3	0.2 ± 0.0	0.2 ± 0.1	1.7 ± 1.0	0.4 ± 0.1	0.5 ± 0.2	0.7 ± 0.3			
Heart	0.8 ± 0.4	0.2 ± 0.1	0.2 ± 0.1	1.4 ± 0.6	0.5 ± 0.2	0.4 ± 0.1	0.5 ± 0.1			
Lung	1.7 ± 0.7	0.4 ± 0.1	0.9 ± 1.0	1.9 ± 0.7	1.0 ± 0.2	0.7 ± 0.1	0.8 ± 0.1			
Tongue	nd	nd	0.3 ± 0.0	2.9 ± 1.2	nd	Nd	nd			

nd, not determined.

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