

Figure S1. Images comparing lack of substrate degradation when fixed eosinophils were added to fibrinogen substrate for 4 hours (left) with substrate degradation from live eosinophils (right)

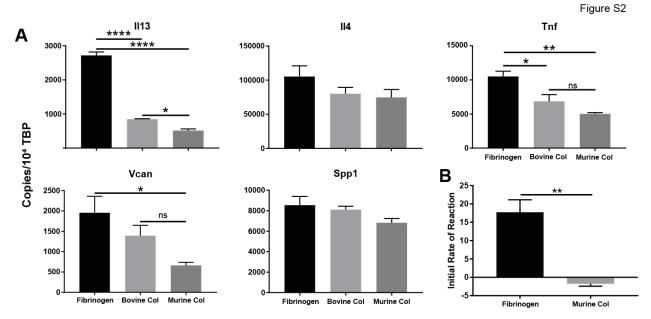


Figure S2. A) Bar graphs showing gene expression comparison between bovine and murine collagen relative to murine fibrinogen. *p<0.5, **p<0.01, ***p<0.001, ****p<0.0001 by one-way ANOVA. B) Biolog metabolic assay comparing metabolic activity of eosinophils exposed to murine fibrinogen vs eosinophils exposed to murine collagen. **p<0.01 by t-test.

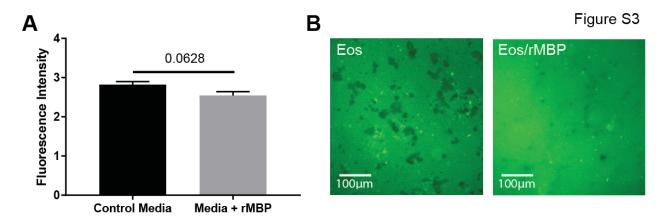


Figure S3. A) Bar graph showing fluorescence intensity of FITC-linked fibrinogen exposed only to cell media control vs. cell media plus 1 μ g/mL rMBP. p=0.0628 by t-test, n=4. B) Visualization of addition of eosinophils with no treatment (left) as compared to those treated with 1 μ g/mL rMBP (right). Images representative of 3 replicates, from which at least 4 fields were imaged.

Mass spectrometry data comparing fibrinogen background control (no eosinophils added) to fibrinogen plus eosinophils										
							Fibrinogen only control (no eosinophils added)		Fibrinogen plus eosinophils	
Protein IDs	Protein names	Gene names	Peptides	Unique peptides	Sequence coverage [%]	Mol. weight [kDa]	LFQ intensity NoEos	LFQ intensity NoEos	LFQ intensity Eos	LFQ intensity Eos
COHKE9;COHKE8;COHKE7;COHKE6;	Histone H2A type 1-H;Histone H2.	Hist1h2ah;H2afj;Hist2h2ac;Hist1h	2	2	21.5	14.135	0	0	0	37034000
E9PV24	Fibrinogen alpha chain; Fibrinopep	Fga	4	4	7.9	87.428	70775000	67089000	60481000	87700000
P18760;P45591	Cofilin-1;Cofilin-2	Cfl1;Cfl2	2	2	23.5	18.559	0	0	27905000	42121000
P49290	Eosinophil peroxidase;Eosinophil p	Epx	4	4	6.6	81.379	0	0	81637000	95396000
P63260;P60710;P63268;P68134;P	Actin, cytoplasmic 2;Actin, cytopla	Actg1;Actb;Actg2;Acta1;Actc1;Act	4	4	14.7	41.792	74386000	102690000	285130000	718610000
Q61878	Bone marrow proteoglycan; Eosine	Prg2	2	2	14.3	24.255	0	0	61286000	84531000
Q8K0E8;CONP02676	Fibrinogen beta chain; Fibrinopept	Fgb	3	3	10.2	54.752	0	0	31581000	42767000
Q8VCM7	Fibrinogen gamma chain	Fgg	2	2	7.1	49.391	24179000	25068000	28231000	36111000

Table \$1. Mass spectrometry data comparing fibrinogen background control (no eosinophilis added) to fibrinogen plus eosinophilis. LFQ = label free quantification, NoEos = fibrinogen only background control, Eos = fibrinogen plus eosinophilis.