

SUPPLEMENTARY MATERIAL

Table S1. Statistics related to Figure 2. Top value is the standard error, SE; bottom is the coefficient of determination, R^2 , for the curve fit from which the inflection point was calculated to derive the K_D . No statistics could be calculated for experiments that did not yield a binding curve and these are left blank.

SARS-CoV-2	Serum Albumin	Cardio-lipin	β 2GPI	Platelet Factor 4	Factor II	Factor VIII	Factor IX	vWF	PDE II	Coll 1
S1				0.015 0.9540				0.016 0.9893	0.032 0.9951	0.047 0.9968
S2										
RBD										
Envelope										0.038 0.9977
Matrix				0.009 0.9940						0.035 0.9965
Nucleoprotein					0.039 0.9919		0.047 0.9893			0.043 0.9964
Other Viruses										
Adenovirus			0.059 0.9829	0.010 0.9820				0.012 0.9933	0.031 0.9614	
Influenza A										
Coxsackie B										
HSV1						0.013 0.9864			0.010 0.9820	0.067 0.7232

Table S2. Statistics related to Figure 6. Top value is the standard error, SE; bottom is the coefficient of determination, R^2 , for the curve fit from which the inflection point was calculated to derive the K_D . No statistics could be calculated for experiments that did not yield a binding curve and these are left blank.

Bacteria	Serum Albumin	Cardio-lipin	β 2GPI	Platelet Factor 4	Factor II	Factor VIII	Factor IX	vWF	PDE II	Coll 1
Staphylococcus		0.038 0.9769	0.083 0.9324	0.008 0.9974					0.011 0.9944	
GAS goat		0.028 0.9529		0.009 0.9940	0.021 0.8405	0.053 0.9584		0.007 0.9965	0.012 0.9850	
GAS rabbit						0.285 0.7923				
Klebsiella		0.130 0.9749	0.046 0.9763						0.014 0.9368	
E. coli		0.019 0.9837	0.030 0.9479	0.003 0.9985					0.009 0.9915	
Clostridium				0.023 0.6745	0.01 0.9915					
M. tuberculosis				0.004 0.9929	0.032 0.9911					

Table S3. Statistics related to Figure 11. Top value is the standard error, SE; bottom is the coefficient of determination, R^2 , for the curve fit from which the inflection point was calculated to derive the K_D . No statistics could be calculated for experiments that did not yield a binding curve and these are left blank.

SARS-CoV-2	Strep Gt	GAS RabHRP	Staph Gt HRP	Kleb Gt HRP	E. coli Gt	Clost HRP Rabbit	M. tb GP
S1	0.010 0.9793	0.01 0.5883	0.002 0.9392				
S2	0.008 0.9925	0.014 0.8170	0.009 0.9944	0.04 0.8917			
RBD	0.015 0.9435	0.005 0.8515					
Envelope							
Matrix							
Nucleocapsid							
Adenovirus Gt							
Influenza A							
CVB blend	0.011 0.9889		0.002 0.9392		0.016 0.9746	0.006 0.9748	
HSV1			0.017 0.9208				