Table 1. Results of pair-wise binding of the anti-HEWL VHHs and binding to HEWL in competition with NAG_3 and Biebrich Scarlet (BS).*

	Group 1						Group 2	
	Epitope 1			Epitope 2			Epitope 3	
	D2-L19	D2-L29	cAbLys2	cAbLys3	D3-L11	D2-L31	D2-L24	D2-L27
Pair-wise binding (%).								
D2-L19	-	7	7	8	12	11	99	100
D2-L29	6	-	9	11	16	0	72	100
cAbLys2	10	10	-	11	17	13	99	100
cAbLys3	11	11	6	-	7	5	100	100
D3-L11	17	18	21	17	-	9	100	99
D2-L31	20	16	18	13	13	-	100	100
D2-L24†	70	75	91	82	78	74	-	5
D2-L27†	86	59	55	71	62	47	0	-
Binding to HEWL in competition with NAG_3 and Biebrich Scarlet (B.S.) (%).								
NAG_3†	95	100	100	13	16	0	99	93
B.S.†	8	10	60	71	3	55	36	93

* The rows represent the first binder, NAG_3 or B.S. and the columns represent the second binder. Cells are shaded according to residual binding of the second binder (5% bins) as determined by surface plasmon resonance on a BIACORE 3000™ instrument. † The values in these rows were obtained by surface plasmon resonance using an IAsys™ instrument. HEWL, hen egg white lysozyme; VHH, variable domain of a heavy chain antibody.