

Table S1 | **Key human antibody Fc region X-ray crystal structures**

Antibody Fc region	Domains crystallized	Fragment details	Complex status	References
IgG1 Fc	C γ 2 and C γ 3	Thr223–Lys447	Uncomplexed	1
			Complexed with fragment B of protein A	2
			Uncomplexed and complexed with Fc γ R1II	38
	C γ 2 and C γ 3	Cys226–Lys447	Complexed with Fc γ R1II	40
	Entire antibody (Dob)	Intact antibody lacking hinge region	Uncomplexed	96
IgE Fc	C ϵ 3 and C ϵ 4 with C ϵ 2–C ϵ 3 linker	Asp330–Lys547 with preceding Cys residue	Uncomplexed	4
	C ϵ 2, C ϵ 3 and C ϵ 4	Lys234–Lys547 with point mutations (Asn265Gln and Asn371Gln) to facilitate crystallization	Complexed with Fc ϵ RI	3
			Uncomplexed	5
IgA Fc	C α 2 and C α 3 lacking tailpiece	Cys242–Lys454	Complexed with Fc α RI	6

Table S2 | **Contributions to Fc stability**

Antibody	Surface area buried in protein–protein domain interactions (\AA^2) per Fc			Inter-heavy-chain disulphide bonds		Fc carbohydrate attachment sites analysed	Domain surface area buried by interaction with carbohydrate (\AA^2) per Fc	
	C γ 2–C γ 2	C γ 2–C γ 3	C γ 3–C γ 3	Hinge	Fc		Penultimate domain	C-terminal domain
IgG	None	1556	2180	2 or more (varies with subclass)	None	Asn297* in C γ 2	C γ 2: 1044	C γ 3: None
IgA	C α 2–C α 2 275	C α 2–C α 3 1662	C α 3–C α 3 2061	None	Three in C α 2: Cys241–Cys241 Cys242–Cys299* x 2	Asn263 in C α 2 Asn459 in tailpiece	C α 2: 930 Unknown	C α 3: 914 Unknown
IgE	C ϵ 3–C ϵ 3 None	C ϵ 3–C ϵ 4 1744	C ϵ 4–C ϵ 4 1860–2326	No hinge present	Two in C ϵ 2: Cys241–Cys328 x 2	Asn394* in C ϵ 3	C ϵ 3: 900	C ϵ 4: None

*Homologous residues.