

**TABLE 8.1** Binding energies and corresponding dissociation constants of different protein-ligand complexes.

Protein	Ligand	Binding Energy (kcal/mol)	Dissociation Constant ( <i>K<sub>d</sub></i> ) (Molar)
Antibody	Antigen	-6 to -11 [54] <sup>*a</sup>	$\sim 10^{-2}$ to $10^{-8}$
Receptor	Hormone	$\sim -12$ [42]	$\sim 10^{-9}$
Enzyme	Substrate	-4 to -8 [55]	$\sim 10^{-3}$ to $10^{-6}$
	Cofactor	-5.5 to -9.5 [42]	$\sim 10^{-4}$ to $10^{-7}$
	Inhibitor	-10 to -15 [54] <sup>*b</sup>	$\sim 10^{-7}$ to $10^{-11}$
	Transition state	-16 to -27 [54] <sup>*c</sup>	$\sim 10^{-12}$ to $10^{-20}$

<sup>\*a</sup>A higher value ( $-14$  kcal/mol ( $K_d \approx 10^{-10}$ )) was obtained for antibodies that underwent a process called ‘*maturation*’. In this process, B-cell lymphocytes adapt to the invading pathogen by producing antibodies with better compatibility with the antigens produced by the pathogen.

<sup>\*b</sup>A similar, yet slightly higher value of  $\approx -18$  kcal/mol ( $K_d = 60$  fM) was measured for the binding of trypsin to pancreatic trypsin inhibitor, which is itself a protein [47].

<sup>\*c</sup>Higher values have also been suggested. For example, Schramm [56] indicated an enzyme-transition state dissociation constant ( $K_d$ ) ranging from  $10^{-14}$  to  $10^{-23}$  mol L<sup>-1</sup>. This corresponds roughly to a binding free energy of  $-19$  to  $-31$  kcal/mol.