## **CH3F5** Molecular Interactions Workshop Additional Material

**Part 1** Peptides are known to assemble into β-sheet structures (below). Estimate how many amino acids would be needed to achieve an association constant of  $10^5 \,\mathrm{M}^{-1}$  between **two** polypeptides aligned in this fashion in chloroform.

HB donor parameter  $\alpha$  amide = 2.9

HB acceptor parameter  $\beta$  amide = 8.3

HB donor parameter  $\alpha$  chloroform = 2.2

HB acceptor parameter  $\beta$  chloroform = 0.8

Part 2 If the corresponding HB acceptor value for DMSO, a competitive solvent is:  $\beta$  = 8.9, estimate what percentage of this solvent would you have to add to the above peptide in chloroform required to reduce the association constant 10,000-fold?

## Part 1 suggested answer:

$$\Delta G = -(\alpha - \alpha_s)(\beta - \beta_s) + 6 \text{ kJ mol}^{-1}$$

For two side-by-side peptide chains, only every OTHER amino acid actually bonds with an ADJACENT peptide. So for n amide HBs, there will be 2n amino acids required, but this multiplier will be used after calculating n.

$$\Delta G = -n(2.9-2.2)(8.3-0.8) + 6 \text{ kJ mol}^{-1}$$

Now  $\Delta G = -RT \ln K_a$ 

If  $R = 0.008314 \text{ kJK}^{-1} \text{mol}^{-1}$ 

Hence  $-n(2.9-2.2)(8.3-0.8) + 6 = -RT \ln 100 000$ 

## Rearrange:

$$-n(2.9-2.2)(8.3-0.8) = -RT \ln 100 000 - 6$$

$$-n(2.9-2.2)(8.3-0.8) = -(RT \ln 100\ 000+6)$$

$$n = (RT \ln 100\ 000+6)/(2.9-2.2)(8.3-0.8)$$

$$n = 34.52/5.25 = 6.6$$

Hence n = 7 hydrogen bonds, which equates to 7 amino acids by rounding up since only every *other* amino acid will hydrogen bond to a given adjacent peptide chain (the  $7^{th}$  hydrogen bond is on the  $7^{th}$  amino acid ... try drawing the structure!)

Part 2 One way of looking at this problem might be ...

$$\Delta G = -RT \ln K_a$$

Hence  $\Delta G = \Delta \Delta G_{HB \text{ Tot}} = -RT \ln 10 \text{ if the } K_a \text{ reduced from } 100 000 \text{ to } 10 \text{ M}^{-1}.$ 

Where 
$$\Delta\Delta G_{HB Tot} = (1-x)\Delta\Delta G_{HB} (CHCl_3) + x\Delta\Delta G_{HB} (DMSO) + 6 \text{ kJ mol}^{-1}$$

Now for the above 7 amino acids (which actually form 6 or 8 hydrogen bonds with an adjacent chain, depending on which you consider):

in CHCl<sub>3</sub>, 
$$\Delta\Delta G_{HB}$$
 (CHCl<sub>3</sub>) =  $-8(2.9-2.2)(8.3-0.8)$  kJ mol<sup>-1</sup>

And for 8 HB in DMSO, 
$$\Delta\Delta G_{HB}$$
 (DMSO) =  $-8(2.9-0.8)(8.3-8.9)$  kJ mol<sup>-1</sup>

Hence  $-(8(2.9-2.2)(8.3-0.8)(1-x) - 8((2.9-0.8)(8.3-8.9)x + 6 = -0.008314 \times 298 \times 10^{-1} \times 10^$ 

$$-42(1-x) +10.08 x + 6 = -0.008314 \times 298 \ln 10$$

$$42x + 10.08 \times -36 = -0.008314 \times 298 \ln 10$$

$$52.08 x = -0.008314 \times 298 \ln 10 + 36$$

=30.295/52.08

$$x = 0.58$$

x = 58 mol% of DMSO required to reduce the association constant from 100 000 to 10 for two 7-mer peptides associated as  $\beta$ -sheet.