

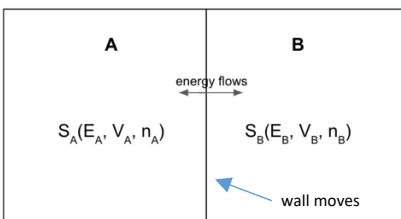
## Hour Exam 2

Useful numbers and formulas you may need are given at the end.

**You must turn in your answer by 10:55 to get full credit.**

- (10 pts) Consider a bag containing three 1-dollar coins and six quarters. You are allowed to pull out exactly one coin in every draw.
  - (1+1) If randomly chosen, **what is** the probability of pulling a single quarter? **Of pulling a** single 1-dollar coin?
  - (2) **What is** the probability of pulling a quarter, a 1-dollar coin, and then another quarter in that order in three draws?
  - (2+1) **What is** the average value  $E$  of a single draw? **Is it possible** to pull out that amount in one draw?
  - (2+1) Let's say we add a half dollar to the bag. **What is** the new average value of a single draw? **Is it possible** to pull out that amount in a single draw?

- (10 pts) Consider an isolated system made up of two subsystems A and B divided by a moveable wall that also allows heat, but not particles, to pass through ( $dn_A=dn_B=0$ ). We allow the subsystems to come to equilibrium:



- (1+1) **Can** work be exchanged between A and B? **Is** entropy extensive (additive)?
  - (1+1) Based on part a., **what is** the change in total entropy  $dS_{tot}$  in terms of  $dS_A$  and  $dS_B$ ? In equilibrium  $dS_{tot}=0$ , therefore **give** the simple relation between  $dS_A$  and  $dS_B$  when A and B are in equilibrium.
  - (1+1+1) **Write down** the general formula for  $dE$  when  $dn=0$  [see 'Useful Information']. Solve for  $dS$ , and **write down** the **two** formulas for  $dS_A$  and  $dS_B$ , putting subscripts "A" or "B" on all variables to distinguish the two subsystems.
  - (1+1+1) **Insert** the two formulas from part c. into part b. at equilibrium. **Use** conservation of total energy  $dE = dE_A + dE_B = 0$  and total volume  $dV = dV_A + dV_B = 0$  to eliminate  $dE_B$  and  $dV_B$  from your formula. **Regroup** your formula to show that in equilibrium, since  $dE_A$  and  $dV_A$  are independent changes, therefore  $T_A = T_B$  and  $P_A = P_B$ .
- (10 pts)  $E$  is a function of  $S$ ,  $V$ , and  $n$ . Therefore  $dE = TdS - PdV + \mu dn$ .  $F$  is a function of  $T$ ,  $V$ , and  $n$ . Let's figure out what  $dF$  is equal to.
    - (2+1) Based on the above, **what** thermodynamic variable is  $\frac{\partial E}{\partial S}$  equal to? **Is it** intensive or extensive?

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b. (2) If  $F$  is the intercept of  $E$  as a function of slope  $\frac{\partial E}{\partial S}$ , **write**  $F$  in terms of  $E$  and simple extensive and intensive variables using your result in a. [Hint: you could look at ‘Useful Information’.]

c. (2) **Write down** the total differential  $dF = \dots$  of your expression for  $F$  in part b., keeping in mind that in general the total differential of a product of two variables,  $AB$ , equals  $d(AB) = AdB + BdA$ .

d. (2+1) Substitute  $dE$  by  $TdS - PdV + \mu dn$  into your result from c. and cancel terms to **write down** the formula for  $dF$ . **What is** the derivative  $\frac{\partial F}{\partial T}$  equal to in terms of a simple variable?

4. (10 pts) Let us calculate the partition function  $Z(T)$  for folding of a peptide with  $N=10$  amino acids. Each amino acid has  $W_{AA}=3$  microstates.

a. (2+1) How many total microstates  $W_{tot}$  does the whole peptide with  $N$  amino acids have? [Hint:  $W$  is a multiplicative quantity, not extensive.] Give a **formula** in terms of  $N$  and  $W_{AA}$ , and **numerical** value.

b. (1) The folded state has  $W_F=1$  microstate. All the other microstates make up the unfolded macrostate. Given your answer in a., **what is**  $W_U$ , the number of microstates in the unfolded state?

c. (2) Assume the folded state has energy  $E_F=0$ , and the unfolded state has energy  $E_U=29$  kJ/mole. **Write down** the canonical partition function  $Z(T)$  for the peptide in terms of  $W_F$ ,  $W_U$ ,  $E_U$ , and  $T$ .

d. (1+1+1+1) **Calculate** the partition function at  $T=298$  K. **Is** the peptide mostly folded or mostly unfolded? **Calculate** the partition function at  $T=350$  K. **Is** the peptide mostly folded or mostly unfolded? [Hint: the partition function  $Z$  is the number of microstates accessible to the system at temperature  $T$ , and the folded state accounts for 1 microstate.]

Useful information:

**Constants:** 1 atomic mass unit  $\approx 1.66 \times 10^{-27}$  kg; mass of electron  $m_e \approx 9.109 \times 10^{-31}$  kg; Planck’s constant  $h \approx 6.626 \times 10^{-34}$  J·s;  $\hbar = h/2\pi$ ; Avogadro’s number  $N_A = 6.02214076 \cdot 10^{23}$ ;  $k_B \approx 1.38 \cdot 10^{-23}$  J/K; gas constant  $R \approx 8.31$  J/mole/K, or  $R \approx -0.08205$  L · atm · mol<sup>-1</sup> · K<sup>-1</sup>; Faraday’s number  $\approx 96485$  Coulombs/mole.  $c \approx 3 \cdot 10^8$  m/s;  $e \approx 1.6 \cdot 10^{-19}$  Coulombs; Avogadro’s number  $N_A = 6.02214076 \cdot 10^{23}$ ; gas constant  $R \approx 8.31$  J/mole/K =  $k_B N_A$ ;

**Partition functions and thermodynamic potentials:**  $\rho_j = 1/W_j$  (constant energy);

$\rho_j = W_j \exp(-E_j/RT)/Z$ ;  $Z = \exp(-F/RT) = \sum W_j \exp(-E_j/RT)$  (constant temperature); average  $A = \sum \rho_j A_j$ ;

$dE = TdS - PdV + \mu dn$ , and this can be solved for  $dS$ ,  $dV$  or  $dn$ .

$E(S,V,\dots)$ ;  $F(T,V,\dots)=E-TS$ ;  $H(S,P,\dots)=E+PV$ ;  $G(T,P,\dots)=H-TS$  all contain the same information.

**Conversions:**  $1 \text{ \AA} = 0.1 \text{ nm} = 100 \text{ pm} = 10^{-10} \text{ m}$ ; wavenumber ( $\text{cm}^{-1}$ ) =  $\frac{10^7}{\lambda \text{ (nm)}}$

**Math:**  $N! = N(N-1)(N-2) \dots 3 \cdot 2 \cdot 1 \approx N^N \cdot N \ln N$ ;  $0! = 1$

**Energy levels:**  $E(n) = -Ry/n^2$ , where  $Ry = 2.1798741 \cdot 10^{-18}$  J (H atom);  $E_n = h\nu(n+1/2)$  (spring)

**Color and wavelength:**

