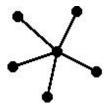
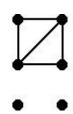
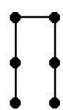
## Probability Models, HW2, due February 7

- **1.** Consider the yeast network which has N=6,018 and  $\alpha=0.0087$ , let K=10 and assume p=0.1.
- (a) Compute the (approximate) speciation probability in our model.
- (b) Compute the speciation probability in Orr's model.
- (c) Roughly how many substitutions do we need in order to get a speciation probability as high as Orr's? You can use an approximation for  $\binom{K}{2}$ .
- 2. For each of the three networks below, let X be the number of interactions (edges) when we choose K=3 nodes. Note that all networks have N=6 and  $N_E=5$ .
- (a) For each network, find P(X = j), j = 0, 1, 2, 3 (easier to use brute force than trying sophisticated combinatorics).
- (b) For each network, compute E[X] both using the definition and the probabilities in (a) and the expression  $\alpha\binom{K}{2}$  (and make sure they coincide).
- (c) For each network, compute the speciation probability for p = 0.1, both exactly using (a) and with our approximation formula.







- **3.** Consider the disjoint network with N nodes,  $N_E = N/2$  edges, choose K nodes at random, and let X be the number of edges in our subgraph. Consider the probability P(X = 0).
- (a) What is P(X = 0) if  $K > N_E$ ?
- (b) What is P(X=0) if N=6 and K=2?
- (c) Find a general expression for P(X = 0) if  $K \leq N_E$ . Remember that there are  $\binom{N}{K}$  ways to choose K nodes out of N so you need to figure out how many of these avoid getting any edges.