a) 1JWA (ATP binding success - area under P/R = .77)

b) 1XEX (ATP binding failure - area under P/R = .05)

c) 3AGC (GLC binding best - area under P/R = .34)

d) 1ACO (GLC binding failure - area under P/R = .01)

e) 3L3X (DHT binding success - area under P/R = .85)
   (protein is semi-transparent to show internal pocket)

f) 1DHT (DHT binding worst - area under P/R = .28)
   (protein is “sliced” to show internal pocket)

g) 1CC5 (HEM binding success - area under P/R = .9)

h) 2Z6F (HEM binding worst - area under P/R = .54)