```
[13] // -- Begin Antimony block converted from wolf2001.xml
// Created by libAntimony v2.9.1
model *wolf2001()
     // Compartments and Species:
     compartment c0, c1, c2;
     species $sul_ex in c0, $eth_ex in c0, $oxy_ex in c0, oxy in c2, $H20 in c2;
     species A3c in c1, aps in c1, $PPi in c1, pap in c1, sul in c1, eth in c1;
     species $A2c in c1, hyd in c1, cys in c1, N2 in c1, $N1 in c1, aco in c1;
     species oah in c1, S1 in c2, $S2 in c2, $C1 in c2, $C2 in c2, $A2m in c2;
     species A3m in c2, $Ho in c1, $Hm in c2;
    // Assignment Rules:
    A2c := Ac - A3c;
    N1 := N - N2;
     S2 := S - S1;
    A2m := Am - A3m;
    // Reactions:
    v1: sul_ex => sul; c0*k_v0/(1 + (cys/Kc)^n);
    v13: $eth_ex => eth; c0*k_v13;
     v2: sul + A3c => aps + $PPi; c1*k2*sul*A3c;
    v10: $oxy_ex => oxy; c0*k_v10;
     v14: oxy => $oxy_ex; c2*k14*oxy;
    v3: aps + A3c => pap + $A2c; c1*k3*aps*A3c;
     v4: pap + 3N2 => hyd + 3$N1; c1*k4*pap*N2;
    v5: hyd + oah => cys; c1*k5*hyd*oah;
     v6: cys => ; c1*k6*cys;
    v7: eth + 2$N1 => aco + 2N2; c1*k7*eth*N1;
    v15: aco => oah; c1*k15*aco;
    v17: hyd => ; c1*k17*hyd;
    v18: oah => ; c1*k18*oah;
    v8: $S2 + aco => S1; c2*k8*aco*S2;
    v9: S1 + 4$N1 => $S2 + 4N2; c2*k9*S1*N1;
    v11a: C1 + Hm + N2 = C2 + Ho + N1; C2*k11*N2*oxy/((a*N2 + oxy)*(1 + C2*k11*N2*oxy/((a*N2 + oxy)*(a*N2 + oxy)*((a*N2 + oxy)*(a*N2 + oxy)*((a*N2 
 (hyd/Kh)^m));
    v11a2: C2 + oxy => C1 + H20; c2*k11*N2*oxy/((a*N2 + oxy)*(1 + (hyd/Kh)^m));
     v16: $A2c + A3m => $A2m + A3c; c2*k16*A3m*A2c;
    v11b: $Ho + $A2m => $Hm + A3m; (c2*3*k11*N2*oxy/((a*N2 + oxy)*(1 + oxy)*(1
 (hyd/Kh)^m))*A2m/(Ka + A2m);
    vLEAK: $Ho => $Hm; 0;
    v12: A3c => $A2c; c1*k12*A3c;
     // Species initializations:
    sul_ex = 0;
    eth_ex = 0;
    oxy_ex = 0;
    oxy = 7;
    oxy has substance_per_volume;
    H20 = 0;
    A3c = 1.5;
    A3c has substance_per_volume;
     aps = 0.5;
     aps has substance_per_volume;
     PPi = 0;
     pap = 0.4;
     pap has substance_per_volume;
     sul = 0.4;
     sul has substance_per_volume;
     eth = 4;
     eth has substance_per_volume;
     A2c has substance_per_volume;
     hyd = 0.5;
     hyd has substance_per_volume;
     cys = 0.3;
     cys has substance_per_volume;
     N2 = 2;
     N2 has substance_per_volume;
     N1 has substance_per_volume;
     aco = 0.3;
     aco has substance_per_volume;
     oah = 1.5;
     oah has substance_per_volume;
     S1 = 1.5;
    S1 has substance_per_volume;
     S2 has substance_per_volume;
    C1 = 0;
    C2 = 0;
    A2m has substance_per_volume;
     A3m = 1.5;
    A3m has substance_per_volume;
    Ho = 0;
    Hm = 0;
    // Compartment initializations:
     c0 = 1;
    c1 = 1;
     c2 = 1;
     // Variable initializations:
    Ac = 2;
    N = 2;
     S = 2;
    Am = 2;
     k_v0 = 1.6;
     Kc = 0.1;
     n = 4;
     k_v13 = 4;
     k2 = 0.2;
     k_v10 = 80;
     k14 = 10;
     k3 = 0.2;
     k4 = 0.2;
     k5 = 0.1;
     k6 = 0.12;
     k7 = 10;
     k15 = 5;
     k17 = 0.02;
     k18 = 1;
     k8 = 10;
     k9 = 10;
     k11 = 10;
     a = 0.1;
     Kh = 0.5;
     m = 4;
     k16 = 10;
     Ka = 1;
     k12 = 5;
     // Other declarations:
     const c0, c1, c2, Ac, N, S, Am, k_v0, Kc, n, k_v13, k2, k_v10, k14, k3;
     const k4, k5, k6, k7, k15, k17, k18, k8, k9, k11, a, Kh, m, k16, Ka, k12;
     // Unit definitions:
    unit substance_per_volume = mole / litre;
    unit substance = mole;
    // Display Names:
    c0 is "external";
     c1 is "cytosol";
    c2 is "mitochondria";
     sul_ex is "SO4_ex";
     eth_ex is "EtOH_ex";
    oxy_ex is "02_ex";
    oxy is "02";
    A3c is "ATP";
     aps is "APS";
     pap is "PAPS";
    sul is "SO4";
     eth is "EtOH";
    A2c is "ADP";
    hyd is "H2S";
    cys is "CYS";
    N2 is "NADH";
    N1 is "NAD";
    aco is "AcCoA";
    oah is "OAH";
    A2m is "ADP_mit";
    A3m is "ATP_mit";
    v11a is "vET1";
    v11a2 is "vET2";
    v11b is "vSYNT";
 end
 // -- End Antimony block
// -- Begin PhraSEDML block converted from main.xml
 // Created by libphrasedml v1.0.5
// Models
 model_m_1 = model "wolf2001" with m=1
 model_m_2 = model "wolf2001" with m=2
 model_m_4 = model "wolf2001" with m=4
 model_m_8 = model "wolf2001" with m=8
// Simulations
sim1 = simulate uniform(0, 100, 5000)
sim2 = simulate uniform(0, 200, 5000)
// Tasks
 task1_100 = run sim1 on model_m_1
 task2_100 = run sim1 on model_m_2
 task4_100 = run sim1 on model_m_4
 task8_100 = run sim1 on model_m_8
 task1_200 = run sim2 on model_m_1
 task2_200 = run sim2 on model_m_2
 task4_200 = run sim2 on model_m_4
 task8_200 = run sim2 on model_m_8
 // Outputs
 plot "Oxygen (100 min)" task1_100.time vs task1_100.oxy, task2_100.oxy, task4_100.oxy,
 task8_100.oxy
 plot "Oxygen (200 min)" task1_200.time vs task1_200.oxy, task2_200.oxy, task4_200.oxy,
 task8_200.oxy
 // -- End PhraSEDML block
              6.5
```



50

20

8

7.5

6.5

40

60

Oxygen (200 min)

100

80

150

100

200

task1\_200.oxy task2\_200.oxy task4\_200.oxy task8\_200.oxy