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%%cardiac cell model
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%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%  
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```
function output=model(t,X,flag_ode)
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```
CL = 300;%ms
```

```
celltype=1; %endo = 0, epi = 1, M = 2
```

```
%extracellular ionic concentrations
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```
nao=140.0;
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```
cao=1.8;
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```
ko=5.4;
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```
%physical constants
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```
R=8314.0;
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```
T=310.0;
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```
F=96485.0;
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```
%cell geometry
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```
L=0.01;
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```
rad=0.0011;
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```
vcell=1000*3.14*rad*rad*L;
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```
Ageo=2*3.14*rad*rad+2*3.14*rad*L;
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```
Acap=2*Ageo;
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```
vmyo=0.68*vcell;
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```
vnsr=0.0552*vcell;
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vjsr=0.0048*vcell;
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```
vss=0.02*vcell;
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```
%parameters setting in control (used in SERCA three-state model)
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```
ADP = 0.015;
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```
Pi = 0;
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```
ATP_in=9.8;
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```
Hi=10^(-4);
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```
%give names to the state vector values
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v=X(1);
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```
nai=X(2);
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```
nass=X(3);
```

```
ki=X(4);
```

```

kss=X(5);
cai=X(6);
cass=X(7);
cansr=X(8);
cajsr=X(9);
m=X(10);
hf=X(11);
hs=X(12);
j=X(13);
hsp=X(14);
jp=X(15);
mL=X(16);
hL=X(17);
hLp=X(18);
a=X(19);
iF=X(20);
iS=X(21);
ap=X(22);
iFp=X(23);
iSp=X(24);
d=X(25);
ff=X(26);
fs=X(27);
fcaf=X(28);
fcas=X(29);
jca=X(30);
nca=X(31);
ffp=X(32);
fcafp=X(33);
xrf=X(34);
xrs=X(35);
xs1=X(36);
xs2=X(37);
xk1=X(38);
JreInp=X(39);
Jrelp=X(40);
CaMKt=X(41);

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%CaMK constants

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KmCaMK=0.15;

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aCaMK=0.05;

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bCaMK=0.00068;
CaMKo=0.05;
KmCaM=0.0015;
%update CaMK
CaMKb=CaMKo*(1.0-CaMKt)/(1.0+KmCaM/cass);
CaMKa=CaMKb+CaMKt;
dCaMKt=aCaMK*CaMKb*(CaMKb+CaMKt)-bCaMK*CaMKt;

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

%reversal potentials
ENa=(R*T/F)*log(nao/nai);
EK=(R*T/F)*log(ko/ki);
PKNa=0.01833;
EKs=(R*T/F)*log((ko+PKNa*nao)/(ki+PKNa*nai));

%convenient shorthand calculations
vffrt=v*F*(R*T);
vfirt=v*F/(R*T);

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%calculate INa
mss=1.0/(1.0+exp((-v+39.57)/9.871));
tm=1.0/(6.765*exp((v+11.64)/34.77)+8.552*exp(-(v+77.42)/5.955));
dm=(mss-m)/tm;
hss=1.0/(1+exp((v+82.90)/6.086));
thf=1.0/(1.432e-5*exp(-(v+1.196)/6.285)+6.149*exp((v+0.5096)/20.27));
ths=1.0/(0.009794*exp(-(v+17.95)/28.05)+0.3343*exp((v+5.730)/56.66));
Ahf=0.99;
Ahs=1.0-Ahf;
dhf=(hss-hf)/thf;
dhs=(hss-hs)/ths;
h=Ahf*hf+Ahs*hs;
jss=hss;
tj=2.038+1.0/(0.02136*exp(-(v+100.6)/8.281)+0.3052*exp((v+0.9941)/38.45));
dj=(jss-j)/tj;
hssp=1.0/(1+exp((v+89.1)/6.086));
thsp=3.0*ths;
dhsp=(hssp-hsp)/thsp;
hp=Ahf*hf+Ahs*hsp;
tjp=1.46*tj;
djp=(jss-jp)/tjp;
GNa=75;

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fINap=(1.0/(1.0+KmCaMK/CaMKa));
INa=GNa*(v-ENa)*m^3.0*((1.0-fINap)*h*j+fINap*hp*jp);

%calculate INaL
mLss=1.0/(1.0+exp((-v+42.85)/5.264));
tmL=tm;
dmL=(mLss-mL)/tmL;
hLss=1.0/(1.0+exp((v+87.61)/7.488));
thL=200.0;
dhL=(hLss-hL)/thL;
hLssp=1.0/(1.0+exp((v+93.81)/7.488));
thLp=3.0*thL;
dhLp=(hLssp-hLp)/thLp;
GNaL=0.0075;
if celltype==1
    GNaL=GNaL*0.6;
end
fINaLp=(1.0/(1.0+KmCaMK/CaMKa));
INaL=GNaL*(v-ENa)*mL*((1.0-fINaLp)*hL+fINaLp*hLp);

%calculate Ito
ass=1.0/(1.0+exp((-v-14.34)/14.82));
ta=1.0515/(1.0/(1.2089*(1.0+exp(-(v-18.4099)/29.3814)))+3.5/(1.0+exp((v+100.0)/29.3814)));
da=(ass-a)/ta;
iss=1.0/(1.0+exp((v+43.94)/5.711));
if celltype==1
    delta_epi=1.0-(0.95/(1.0+exp((v+70.0)/5.0)));
else
    delta_epi=1.0;
end
tiF=4.562+1/(0.3933*exp((-v+100.0)/100.0)+0.08004*exp((v+50.0)/16.59));
tiS=23.62+1/(0.001416*exp((-v+96.52)/59.05)+1.780e-8*exp((v+114.1)/8.079));
tiF=tiF*delta_epi;
tiS=tiS*delta_epi;
AiF=1.0/(1.0+exp((v-213.6)/151.2));
AiS=1.0-AiF;
diF=(iss-iF)/tiF;
diS=(iss-iS)/tiS;
i=AiF*iF+AiS*iS;
assp=1.0/(1.0+exp((-v-24.34)/14.82));
dap=(assp-ap)/ta;
dti_develop=1.354+1.0e-4/(exp((v-167.4)/15.89)+exp(-(v-12.23)/0.2154));
dti_recover=1.0-0.5/(1.0+exp((v+70.0)/20.0));
tiFp=dti_develop*dti_recover*tiF;

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tiSp=dti_develop*dti_recover*tiS;
diFp=(iss-iFp)/tiFp;
diSp=(iss-iSp)/tiSp;
ip=AiF*iFp+AiS*iSp;
Gto=0.02;
if celltype==1
    Gto=Gto*4.0;
elseif celltype==2
    Gto=Gto*4.0;
end
fltop=(1.0/(1.0+KmCaMK/CaMKa));
Ito=Gto*(v-EK)*((1.0-fltop)*a*i+fltop*ap*ip);

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[vcycle, pp] = Tran_SERCA_3state(cai,cajsr,Hi,ATP_in,ADP,Pi); %adding the SERCA three-
state model in the cell model

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dss=1.0/(1.0+exp((-v+3.940)/4.230));
td=0.6+1.0/(exp(-0.05*(v+6.0))+exp(0.09*(v+14.0)));
dd=(dss-d)/td;
fss=1.0/(1.0+exp((v+19.58)/3.696));
tff=7.0+1.0/(0.0045*exp(-(v+20.0)/10.0)+0.0045*exp((v+20.0)/10.0));
tfs=1000.0+1.0/(0.000035*exp(-(v+5.0)/4.0)+0.000035*exp((v+5.0)/6.0));
Aff=0.6;
Afs=1.0-Aff;
dff=(fss-ff)/tff;
dfs=(fss-fs)/tfs;
f=Aff*ff+Afs*fs;
fcass=fss;
tfcaf=7.0+1.0/(0.04*exp(-(v-4.0)/7.0)+0.04*exp((v-4.0)/7.0));
tfcas=100.0+1.0/(0.00012*exp(-v/3.0)+0.00012*exp(v/7.0));
Afcaf=0.3+0.6/(1.0+exp((v-10.0)/10.0));
Afcas=1.0-Afcaf;
dfcaf=(fcass-fcaf)/tfcaf;
dfcas=(fcass-fcas)/tfcas;
fca=Afcaf*fcaf+Afcas*fcas;
tjca=75.0;
djca=(fcass-jca)/tjca;
tffp=2.5*tff;
dffp=(fss-ffp)/tffp;
fpp=Aff*ffp+Afs*fs;
tfcafpp=2.5*tfcaf;
dfcafpp=(fcass-fcafpp)/tfcafpp;

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fcap=Afcap*fcafp+Afcas*fcas;
Kmn=0.002;
k2n=1000.0;
km2n=jca*1.0;
anca=1.0/(k2n/km2n+(1.0+Kmn/cass)^4.0);
dnca=anca*k2n-nca*km2n;
PhiCaL=4.0*vffrt*(cass*exp(2.0*vfrt)-0.341*cao)/(exp(2.0*vfrt)-1.0);
PhiCaNa=1.0*vffrt*(0.75*nass*exp(1.0*vfrt)-0.75*nao)/(exp(1.0*vfrt)-1.0);
PhiCaK=1.0*vffrt*(0.75*kss*exp(1.0*vfrt)-0.75*ko)/(exp(1.0*vfrt)-1.0);
zca=2.0;
PCa=0.0001;
if celltype==1
    PCa=PCa*1.2;
elseif celltype==2
    PCa=PCa*2.5;
end
PCap=1.1*PCa;
PCaNa=0.00125*PCa;
PCaK=3.574e-4*PCa;
PCaNap=0.00125*PCap;
PCaKp=3.574e-4*PCap;
fICaLp=(1.0/(1.0+KmCaMK/CaMKa));
ICaL=(1.0-fICaLp)*PCa*PhiCaL*d*(f*(1.0-nca)+jca*fca*nca)+fICaLp*PCap*PhiCaL*d*(fp*(1.0-
nca)+jca*fcap*nca);
ICaNa=(1.0-fICaLp)*PCaNa*PhiCaNa*d*(f*(1.0-
nca)+jca*fca*nca)+fICaLp*PCaNap*PhiCaNa*d*(fp*(1.0-nca)+jca*fcap*nca);
ICaK=(1.0-fICaLp)*PCaK*PhiCaK*d*(f*(1.0-
nca)+jca*fca*nca)+fICaLp*PCaKp*PhiCaK*d*(fp*(1.0-nca)+jca*fcap*nca);

%calculate IKr
xrss=1.0/(1.0+exp((-v+8.337)/6.789));
txrf=12.98+1.0/(0.3652*exp((v-31.66)/3.869)+4.123e-5*exp((-v-47.78)/20.38));
txrs=1.865+1.0/(0.06629*exp((v-34.70)/7.355)+1.128e-5*exp((-v-29.74)/25.94));
Axrf=1.0/(1.0+exp((v+54.81)/38.21));
Axrs=1.0-Axrf;
dxrf=(xrss-xrf)/txrf;
dxrs=(xrss-xrs)/txrs;
xr=Axrf*xrf+Axrs*xrs;
rkr=1.0/(1.0+exp((v+55.0)/75.0))*1.0/(1.0+exp((v-10.0)/30.0));
GKr=0.046;
if celltype==1
    GKr=GKr*1.3;
elseif celltype==2
    GKr=GKr*0.8;

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end
IKr=GKr*sqrt(ko/5.4)*xr*rkr*(v-EK);

%calculate IKs
xs1ss=1.0/(1.0+exp((-v+11.60)/8.932));
txs1=817.3+1.0/(2.326e-4*exp((v+48.28)/17.80)+0.001292*exp((-v+210.0)/230.0));
dxs1=(xs1ss-xs1)/txs1;
xs2ss=xs1ss;
txs2=1.0/(0.01*exp((v-50.0)/20.0)+0.0193*exp((-v+66.54)/31.0));
dxs2=(xs2ss-xs2)/txs2;
KsCa=1.0+0.6/(1.0+(3.8e-5/cai)^1.4);
GKs=0.0034;
if celltype==1
    GKs=GKs*1.4;
end
IKs=GKs*KsCa*xs1*xs2*(v-EKs);

xk1ss=1.0/(1.0+exp(-(v+2.5538*ko+144.59)/(1.5692*ko+3.8115)));
txk1=122.2/(exp((-v+127.2)/20.36)+exp((v+236.8)/69.33));
dxk1=(xk1ss-xk1)/txk1;
rk1=1.0/(1.0+exp((v+105.8-2.6*ko)/9.493));
GK1=0.1908;
if celltype==1
    GK1=GK1*1.2;
elseif celltype==2
    GK1=GK1*1.3;
end
IK1=GK1*sqrt(ko)*rk1*xk1*(v-EK);

%calculate INaCa_i
kna1=15.0;
kna2=5.0;
kna3=88.12;
kasymm=12.5;
wna=6.0e4;
wca=6.0e4;
wnaca=5.0e3;
kcaon=1.5e6;
kcaoff=5.0e3;
qna=0.5224;
qca=0.1670;
hca=exp((qca*v*F)/(R*T));
hna=exp((qna*v*F)/(R*T));
h1=1+nai/kna3*(1+hna);

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h2=(nai*hna)/(kna3*h1);
h3=1.0/h1;
h4=1.0+nai/kna1*(1+nai/kna2);
h5=nai*nai/(h4*kna1*kna2);
h6=1.0/h4;
h7=1.0+nao/kna3*(1.0+1.0/hna);
h8=nao/(kna3*hna*h7);
h9=1.0/h7;
h10=kasymm+1.0+nao/kna1*(1.0+nao/kna2);
h11=nao*nao/(h10*kna1*kna2);
h12=1.0/h10;
k1=h12*cao*kcaon;
k2=kcaoff;
k3p=h9*wca;
k3pp=h8*wnaca;
k3=k3p+k3pp;
k4p=h3*wca/hca;
k4pp=h2*wnaca;
k4=k4p+k4pp;
k5=kcaoff;
k6=h6*cai*kcaon;
k7=h5*h2*wna;
k8=h8*h11*wna;
x1=k2*k4*(k7+k6)+k5*k7*(k2+k3);
x2=k1*k7*(k4+k5)+k4*k6*(k1+k8);
x3=k1*k3*(k7+k6)+k8*k6*(k2+k3);
x4=k2*k8*(k4+k5)+k3*k5*(k1+k8);
E1=x1/(x1+x2+x3+x4);
E2=x2/(x1+x2+x3+x4);
E3=x3/(x1+x2+x3+x4);
E4=x4/(x1+x2+x3+x4);
KmCaAct=150.0e-6;
allo=1.0/(1.0+(KmCaAct/cai)^2.0);
zna=1.0;
JncxNa=3.0*(E4*k7-E1*k8)+E3*k4pp-E2*k3pp;
JncxCa=E2*k2-E1*k1;
Gncx=0.0008;
if celltype==1
    Gncx=Gncx*1.1;
elseif celltype==2
    Gncx=Gncx*1.4;
end
INaCa_i=0.8*Gncx*allo*(zna*JncxNa+zca*JncxCa);

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```

%calculate INaCa_ss
h1=1+nass/kna3*(1+hna);
h2=(nass*hna)/(kna3*h1);
h3=1.0/h1;
h4=1.0+nass/kna1*(1+nass/kna2);
h5=nass*nass/(h4*kna1*kna2);
h6=1.0/h4;
h7=1.0+nao/kna3*(1.0+1.0/hna);
h8=nao/(kna3*hna*h7);
h9=1.0/h7;
h10=kasymm+1.0+nao/kna1*(1+nao/kna2);
h11=nao*nao/(h10*kna1*kna2);
h12=1.0/h10;
k1=h12*cao*kcaon;
k2=kcaoff;
k3p=h9*wca;
k3pp=h8*wnaca;
k3=k3p+k3pp;
k4p=h3*wca/hca;
k4pp=h2*wnaca;
k4=k4p+k4pp;
k5=kcaoff;
k6=h6*cass*kcaon;
k7=h5*h2*wna;
k8=h8*h11*wna;
x1=k2*k4*(k7+k6)+k5*k7*(k2+k3);
x2=k1*k7*(k4+k5)+k4*k6*(k1+k8);
x3=k1*k3*(k7+k6)+k8*k6*(k2+k3);
x4=k2*k8*(k4+k5)+k3*k5*(k1+k8);
E1=x1/(x1+x2+x3+x4);
E2=x2/(x1+x2+x3+x4);
E3=x3/(x1+x2+x3+x4);
E4=x4/(x1+x2+x3+x4);
KmCaAct=150.0e-6;
allo=1.0/(1.0+(KmCaAct/cass)^2.0);
JncxNa=3.0*(E4*k7-E1*k8)+E3*k4pp-E2*k3pp;
JncxCa=E2*k2-E1*k1;
INaCa_ss=0.2*Gncx*allo*(zna*JncxNa+zca*JncxCa);

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%calculate INaK

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```

k1p=949.5;
k1m=182.4;
k2p=687.2;
k2m=39.4;

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k3p=1899.0;
k3m=79300.0;
k4p=639.0;
k4m=40.0;
Knai0=9.073;
Knao0=27.78;
delta=-0.1550;
Knai=Knai0*exp((delta*v*F)/(3.0*R*T));
Knao=Knao0*exp(((1.0-delta)*v*F)/(3.0*R*T));
Kki=0.5;
Kko=0.3582;
MgADP=0.05;
MgATP=9.8;
Kmgatp=1.698e-7;
H=1.0e-7;
eP=4.2;
Khp=1.698e-7;
Knap=224.0;
Kxkur=292.0;
P=eP/(1.0+H/Khp+nai/Knap+ki/Kxkur);
a1=(k1p*(nai/Knai)^3.0)/((1.0+nai/Knai)^3.0+(1.0+ki/Kki)^2.0-1.0);
b1=k1m*MgADP;
a2=k2p;
b2=(k2m*(nao/Knao)^3.0)/((1.0+nao/Knao)^3.0+(1.0+ko/Kko)^2.0-1.0);
a3=(k3p*(ko/Kko)^2.0)/((1.0+nao/Knao)^3.0+(1.0+ko/Kko)^2.0-1.0);
b3=(k3m*P*H)/(1.0+MgATP/Kmgatp);
a4=(k4p*MgATP/Kmgatp)/(1.0+MgATP/Kmgatp);
b4=(k4m*(ki/Kki)^2.0)/((1.0+nai/Knai)^3.0+(1.0+ki/Kki)^2.0-1.0);
x1=a4*a1*a2+b2*b4*b3+a2*b4*b3+b3*a1*a2;
x2=b2*b1*b4+a1*a2*a3+a3*b1*b4+a2*a3*b4;
x3=a2*a3*a4+b3*b2*b1+b2*b1*a4+a3*a4*b1;
x4=b4*b3*b2+a3*a4*a1+b2*a4*a1+b3*b2*a1;
E1=x1/(x1+x2+x3+x4);
E2=x2/(x1+x2+x3+x4);
E3=x3/(x1+x2+x3+x4);
E4=x4/(x1+x2+x3+x4);
zk=1.0;
JnakNa=3.0*(E1*a3-E2*b3);
JnakK=2.0*(E4*b1-E3*a1);
Pnak=30;
if celltype==1
    Pnak=Pnak*0.9;
elseif celltype==2
    Pnak=Pnak*0.7;

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end
INaK=Pnak*(zna*JnakNa+zk*JnakK);

%calculate IKb
xkb=1.0/(1.0+exp(-(v-14.48)/18.34));
GKb=0.003;
if celltype==1
    GKb=GKb*0.6;
end
IKb=GKb*xkb*(v-EK);

%calculate INab
PNab=3.75e-10;
INab=PNab*vffrt*(nai*exp(vfirt)-nao)/(exp(vfirt)-1.0);

%calculate ICab
PCab=2.5e-8;
ICab=PCab*4.0*vffrt*(cai*exp(2.0*vfirt)-0.341*cao)/(exp(2.0*vfirt)-1.0);

%calculate IpCa
GpCa=0.0005;
IpCa=GpCa*cai/(0.0005+cai);

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%calculate the stimulus current, Istim
amp=-80.0;
duration=0.5;

if (t-floor(t/CL)*CL <= duration)  && (t-floor(t/CL)*CL >0)
    Istim=amp;
else
    Istim=0;
end

%update the membrane voltage
dv= -
(INa+INaL+Ito+ICaL+ICaNa+ICaK+IKr+IKs+IK1+INaCa_i+INaCa_ss+INaK+INab+IKb+IpCa+
ICab+Istim);

%calculate diffusion fluxes
JdiffNa=(nass-nai)/2.0;
JdiffK=(kss-ki)/2.0;

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Jdiff=(cass-cai)/0.2;

%calculate ryanodine receptor calcium induced calcium release from the jsr
bt=4.75;
a_rel=0.5*bt;
Jrel_inf=a_rel*(-lCaL)/(1.0+(1.5/cajsr)^8.0);
if celltype==2
    Jrel_inf=Jrel_inf*1.7;
end
tau_rel=2*bt/(1.0+0.0123/cajsr);

if tau_rel<0.001
    tau_rel=0.001;
end
dJrelnp=(Jrel_inf-Jrelnp)/tau_rel;
btp=1.25*bt;
a_relp=0.5*btp;
Jrel_infp=a_relp*(-lCaL)/(1.0+(1.5/cajsr)^8.0);
if celltype==2
    Jrel_infp=Jrel_infp*1.7;
end
tau_relp=2*btp/(1.0+0.0123/cajsr);
if tau_relp<0.001
    tau_relp=0.001;
end

dJrelp=(Jrel_infp-Jrelp)/tau_relp;
fJrelp=(1.0/(1.0+KmCaMK/CaMKa));
Jrel=(1.0-fJrelp)*Jrelnp+fJrelp*Jrelp;

%calculate serca pump, ca uptake flux
Jupnp=0.004375*cai/(cai+0.00092);
Jupp=2.75*0.004375*cai/(cai+0.00092-0.00017);
if celltype==1
    Jupnp=Jupnp*1.3;
    Jupp=Jupp*1.3;
end
fJupp=(1.0/(1.0+KmCaMK/CaMKa));
% Jup=(1.0-fJupp)*Jupnp+fJupp*Jupp-Jleak; %%%original equation in Ord model
Jleak=0.0039375*cansr/15.0;
Jup = 0.00179*vcycle-Jleak; %%% calculate Jup by using the cycle rate after adding the SERCA
three-state model
%calculate tranlocation flux
Jtr=(cansr-cajsr)/100.0;

```

```

%calcium buffer constants
cmdnmax=0.05;
if celltype==1
    cmdnmax=cmdnmax*1.3;
end
kmcmdn=0.00238;
trpnmax=0.07;
kmtrpn=0.0005;
BSRmax=0.047;
KmBSR=0.00087;
BSLmax=1.124;
KmBSL=0.0087;
csqnmax=10.0;
kmcsqn=0.8;

%update intracellular concentrations, using buffers for cai, cass, cajsr
dnai=-(INa+INaL+3.0*INaCa_i+3.0*INaK+INab)*Acap/(F*vmyo)+JdiffNa*vss/vmyo;
dnass=-(ICaNa+3.0*INaCa_ss)*Acap/(F*vss)-JdiffNa;

dki=-(Ito+IKr+IKs+IK1+IKb+Istim-2.0*INaK)*Acap/(F*vmyo)+JdiffK*vss/vmyo;
dkss=-(ICaK)*Acap/(F*vss)-JdiffK;

Bcai=1.0/(1.0+cmdnmax*kmcmdn/(kmcmdn+cai)^2.0+trpnmax*kmtrpn/(kmtrpn+cai)^2.0);
dcai=Bcai*(-(IpCa+ICab-2.0*INaCa_i)*Acap/(2.0*F*vmyo)-Jup*vnsr/vmyo+Jdiff*vss/vmyo);

Bcass=1.0/(1.0+BSRmax*KmBSR/(KmBSR+cass)^2.0+BSLmax*KmBSL/(KmBSL+cass)^2.0);
dcass=Bcass*(-(ICaL-2.0*INaCa_ss)*Acap/(2.0*F*vss)+Jrel*vjsr/vss-Jdiff);

dcansr=Jup-Jtr*vjsr/vnsr;

Bcajsr=1.0/(1.0+csqnmax*kmcsqn/(kmcsqn+cajsr)^2.0);
dcajsr=Bcajsr*(Jtr-Jrel);

cajsr_buff = csqnmax*(cajsr/(cajsr+kmcsqn));

%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
output the state vector when ode_flag==1, and the calculated currents and fluxes otherwise
if flag_ode==1
    output=[dv dnai dnass dki dkss dcai dcass dcansr dcajsr dm dhf dhs dj dhsp djp dmL dhL
    dhLp da diF diS dap diFp diSp dd dff dfs dfcaf dfcas djca dnca dffp dfcafp dxrf dxrs dxs1 dxs2
    dxk1 dJrelnp dJrelp dCaMKt]';
else

```

```
output=[INa INaL Ito ICaL IKr IKs IK1 INaCa_i INaCa_ss INaK IKb INab ICab IpCa Jdiff  
JdiffNa JdiffK Jup Jleak Jtr Jrel CaMKa Istim cajsr_buff];  
end
```

%SERCA THREE-STATE MODEL

%%

%% Three-state SERCA model %%%%%%%%%%

%%

% This function outputs the SERCA three-state model cycle rate

% and phosphorylation proportion as a function of Cai, Casr, ATP, ADP and Pi

% All concentrations are in units of mM - including H_i

% Author: Kenneth Tran

% Last updated: 25/09/08

function [v_cycle, phosph] = Tran_SERCA_3state(Ca_i,Ca_sr,H_i,ATP,ADP,Pi)

%Assigning the parameter values

k_p1 = 25900;

k_p2 = 2540;

k_p3 = 20.5;

k_m1 = 2;

k_m2 = 67200;

k_m3 = 149;

kdcai = 0.91;

kdcasr = 2.24;

kdh1 = 1.09e-5;

kdhi = 3.54e-3;

kdhsr = 1.05e-8;

kdh = 7.24e-5;

n_H = 2.0;

T_Cai = Ca_i/kdcai;

T_Casr = Ca_sr/kdcasr;

T_H1 = H_i/kdh1;

T_Hi = (H_i^n_H)/kdhi;

T_Hsr = (H_i.^n_H)/kdhsr;

T_H = H_i/kdh;

% Apparent forward rate constants

a_p1 = k_p1*ATP;

a_p2 = k_p2*T_Cai*T_Cai/(T_Cai^2 + T_Cai^2.*T_Hi + T_Hi*(1+T_H1));

a_p3 = (k_p3*T_Hsr)/(T_Casr*T_Casr*T_H + T_H*(1) + T_Hsr*(1+T_H));

% Apparent backward rate constants

a_m1 = k_m1*T_Hi/(T_Cai^2 + T_Cai^2.*T_Hi + T_Hi*(1+T_H1));

a_m2 = (k_m2*ADP*T_Casr*T_Casr*T_H)/(T_Casr*T_Casr*T_H + T_H + T_Hsr*(1+T_H));

a_m3 = k_m3*Pi;

% Calculating steady state flux using the diagram method

$s1 = a_{p2} \cdot a_{p3} + a_{m1} \cdot a_{p3} + a_{m1} \cdot a_{m2};$

$s2 = a_{p1} \cdot a_{p3} + a_{m2} \cdot a_{p1} + a_{m2} \cdot a_{m3};$

$s3 = a_{p1} \cdot a_{p2} + a_{m3} \cdot a_{m1} + a_{m3} \cdot a_{p2};$

$sum = s1 + s2 + s3;$

% Calculating the cycle rate

$v_{cycle} = (a_{p1} \cdot a_{p2} \cdot a_{p3} - a_{m1} \cdot a_{m2} \cdot a_{m3}) / sum;$

% Proportion of states that are phosphorylated

$phosph = (s3) / (s1 + s2 + s3);$

%SOLVE FUNCTION

%%

%% Solve function %%%%%%%%%%

%%

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% Washington University in St. Louis.

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% DAMAGE.

% MATLAB Implementation of the O'Hara-Rudy dynamic (ORd) model for the
% undiseased human ventricular action potential and calcium transient

%

% The ORd model is described in the article "Simulation of the Undiseased
% Human Cardiac Ventricular Action Potential: Model Formulation and
% Experimental Validation"

% by Thomas O'Hara, Laszlo Virag, Andras Varro, and Yoram Rudy

%
% The article and supplemental materials are freely available in the
% Open Access journal PLoS Computational Biology
% Link to Article:
% <http://www.ploscompbiol.org/article/info:doi/10.1371/journal.pcbi.1002061>
%
% Email: tom.ohara@gmail.com / rudy@wustl.edu
% Web: <http://rudylab.wustl.edu>
% We use the function to solve currents and ion concentrations.

close all

clear all

%initial conditions for state variables

v=-87;
nai=7;
nass=nai;
ki=145;
kss=ki;
cai=1.0e-4;
cass=cai;
cansr=1.2;
cajsr=cansr;
m=0;
hf=1;
hs=1;
j=1;
hsp=1;
jp=1;
mL=0;
hL=1;
hLp=1;
a=0;
iF=1;
iS=1;
ap=0;
iFp=1;
iSp=1;
d=0;
ff=1;
fs=1;
fcaf=1;
fcas=1;
jca=1;

```

nca=0;
ffp=1;
fcafp=1;
xrf=0;
xrs=0;
xs1=0;
xs2=0;
xk1=1;
JreInp=0;
Jrelp=0;
CaMKt=0;

timeStep1=1;
Length_array=100000;
T_array1=zeros(Length_array,1);
T_array1(1) = 500;
options1 = odeiset('MaxStep',0.1);           % Set integrator options

%X0 is the vector for initial sconditions for state variables
X0=[v nai nass ki kss cai cass cansr cajsr m hf hs j hsp jp mL hL hLp a iF iS ap iFp iSp d ff fs
fcaf fcas jca nca ffp fcafp xrf xrs xs1 xs2 xk1 JreInp Jrelp CaMKt]';
CL=300;%pacing cycle length in ms
beats=10;%number of beats in the simulation
options=odeiset('MaxStep',1);%options for ode solver
X = [];
T = [];

for m = [1:beats]
[TT XX]=ode23t(@model,[0 CL],X0',options1,1);
X = [X;XX];
T = [T;CL*(m-1)+TT];
X0 = XX(size(XX,1),:);
m
end

%rename values in the state variables vector
v=X(:,1);
nai=X(:,2);
nass=X(:,3);
ki=X(:,4);
kss=X(:,5);
cai=X(:,6);
cass=X(:,7);

```

```

cansr=X(:,8);
cajsr=X(:,9);
m=X(:,10);
hf=X(:,11);
hs=X(:,12);
j=X(:,13);
hsp=X(:,14);
jp=X(:,15);
mL=X(:,16);
hL=X(:,17);
hLp=X(:,18);
a=X(:,19);
iF=X(:,20);
iS=X(:,21);
ap=X(:,22);
iFp=X(:,23);
iSp=X(:,24);
d=X(:,25);
ff=X(:,26);
fs=X(:,27);
fcaf=X(:,28);
fcas=X(:,29);
jca=X(:,30);
nca=X(:,31);
ffp=X(:,32);
fcafp=X(:,33);
xrf=X(:,34);
xrs=X(:,35);
xs1=X(:,36);
xs2=X(:,37);
xk1=X(:,38);
JreInp=X(:,39);
Jrelp=X(:,40);
CaMKt=X(:,41);

```

```

% calculate and name dependent variables for the final beat in the
% simulation (i.e. currents and fluxes)

```

```

for i=[1:size(X,1)];
    IsJs=model(T(i),X(i,:),0);
    INa(i)=IsJs(1);
    INaL(i)=IsJs(2);
    Ito(i)=IsJs(3);
    ICaL(i)=IsJs(4);
    IKr(i)=IsJs(5);

```

```

IKs(i)=IsJs(6);
IK1(i)=IsJs(7);
INaCa_i(i)=IsJs(8);
INaCa_ss(i)=IsJs(9);
INaK(i)=IsJs(10);
IKb(i)=IsJs(11);
INab(i)=IsJs(12);
ICab(i)=IsJs(13);
IpCa(i)=IsJs(14);
Jdiff(i)=IsJs(15);
JdiffNa(i)=IsJs(16);
JdiffK(i)=IsJs(17);
Jup(i)=IsJs(18);
Jleak(i)=IsJs(19);
Jtr(i)=IsJs(20);
Jrel(i)=IsJs(21);
CaMKa(i)=IsJs(22);
Istim(i)=IsJs(23);
end

%cell geometry
L=0.01;
rad=0.0011;
vcell=1000*3.14*rad*rad*L;
vnssr=0.0552*vcell;
vjsr=0.0048*vcell;
Casr = (vjsr*X(:,8) + vnssr*X(:,9))/(vjsr+vnssr); %Ca2+ concentration in the SR

figure(1)
plot(T,v);
xlabel('T (ms)');
ylabel('V (mv)');
figure(2)
plot(T,X(:,6));
xlabel('T (ms)');
ylabel('Cai (mmol/L)');
figure(3)
plot(T,Casr);
xlabel('T (ms)');
ylabel('Casr (mmol/L)');

```

%figure1.m

%%%

%%%figure1.m %%%%%%%%%%

%%%

clear all;

close all;

clc;

load A11; %Calculated after 1000 beats in ischemic case 2 with CL=300ms; There are instructions about its content in the following

T3 = A11(:,1);

V = A11(:,2);

Cai = A11(:,3);

Casr = A11(:,4);

Jrel = A11(:,5);

Ito = A11(:,6);

ICaL = A11(:,7);

subplot(4,2,1)

plot(T3,V,'k','linewidth',1);

xlabel('T (ms)');

ylabel('V (mv)');

axis([0 3000 -100 50])

subplot(4,2,3)

plot(T3,Cai,'k','linewidth',1);

axis([0 3000 0 0.003])

xlabel('T (ms)');

ylabel('Ca_i (mmol/L)');

subplot(4,2,5)

plot(T3,Casr,'k','linewidth',1);

xlabel('T (ms)', 'FontSize', 9);

ylabel('Ca_s_r (mmol/L)', 'FontSize', 9);

axis([0 3000 2.5 4]);

a=find(T3==300);

b=find(T3==600);

subplot(4,2,7)

plot(T3(1:a(1)),Ito(1:a(1)),'k','linewidth',1);

```

hold on
plot(T3(a(2):b(1))-300,Ito(a(2):b(1)),'r--','linewidth',1);
xlabel('T (ms)');
ylabel('I_t_o (uA/uF)');
axis([-10 300 0 4]);

```

```

subplot(4,2,2)
plot(T3(1:a(1)),V(1:a(1)),'k','linewidth',1);
hold on
plot(T3(a(2):b(1))-300,V(a(2):b(1)),'r--','linewidth',1);
xlabel('T (ms)');
ylabel('V (mv)');
axis([-10 300 -100 50])
axes('position',[0.788714285714285 0.850938095238102 0.098 0.045],'FontSize',8);
plot(T3(12:48),V(12:48),'linewidth',1);
hold on
plot(T3(720:755)-300,V(720:755),'r--','linewidth',1);
axis([5 20 22 32]);

```

```

subplot(4,2,4)
plot(T3(1:a(1)),Cai(1:a(1)),'k','linewidth',1);
hold on
plot(T3(a(2):b(1))-300,Cai(a(2):b(1)),'r--','linewidth',1)
xlabel('T (ms)');
ylabel('Ca_i (mmol/L)');
axis([-10 300 0 0.003])
axes('position',[0.783357142857143 0.633981937602632 0.09800000000000001
0.045],'FontSize',8);
plot(T3(684:710),Cai(684:710),'k','linewidth',1);
hold on
plot(T3(1384:1411)-300,Cai(1384:1411),'r--','linewidth',1);
axis([290 300 0.00037 0.00041]);

```

```

subplot(4,2,6)
plot(T3(1:a(1)),Jrel(1:a(1)),'k','linewidth',1);
hold on
plot(T3(a(2):b(1))-300,Jrel(a(2):b(1)),'r--','linewidth',1)
xlabel('T (ms)');
ylabel('J_r_e_l (mM/ms)');
axis([-10 300 0 2]);

```

```
subplot(4,2,8)
plot(T3(1:a(1)),lCaL(1:a(1)), 'k', 'linewidth', 1);
hold on
plot(T3(a(2):b(1))-300, lCaL(a(2):b(1)), 'r--', 'linewidth', 1)
xlabel('T (ms)');
ylabel('I_C_a_L (uA/uF)');
axis([-10 300 -4 0]);
```

```
annotation('textbox',[0.0423 0.89 0.059 0.067], 'String',{'a'}, 'LineStyle','none');
annotation('textbox',[0.0423 0.67 0.059 0.067], 'String',{'b'}, 'LineStyle','none');
annotation('textbox',[0.0423 0.46 0.059 0.067], 'String',{'c'}, 'LineStyle','none');
annotation('textbox',[0.0423 0.24 0.059 0.067], 'String',{'d'}, 'LineStyle','none');
annotation('textbox',[0.475 0.89 0.059 0.067], 'String',{'e'}, 'LineStyle','none');
annotation('textbox',[0.475 0.67 0.059 0.067], 'String',{'f'}, 'LineStyle','none');
annotation('textbox',[0.475 0.46 0.059 0.067], 'String',{'g'}, 'LineStyle','none');
annotation('textbox',[0.475 0.24 0.059 0.067], 'String',{'h'}, 'LineStyle','none');
```


%figure2.m

%%%

%%%figure2.m %%%%%%%%%%

%%%

clear all;

close all;

clc;

subplot(2,1,1)

load A20; %Its contents are explained in the following

load A21; %Its contents are explained in the following

casr_case1a = A20(:,1); %Recordings of SR calcium ion concentration of each heart beat(the total number is 1000) in case 1 with CL = 250ms;

casr_case2a = A20(:,2); %Recordings of SR calcium ion concentration of each heart beat in case 2 with CL = 250ms;

casr_case3a = A20(:,3); %Recordings of SR calcium ion concentration of each heart beat in case 3 with CL = 250ms;

casr_cona = A20(:,4); %Recordings of SR calcium ion concentration of each heart beat in control condition with CL = 250ms;

carelease_case1a = A20(:,5); %Recordings of JSR released calcium ion concentration of each heart beat in case 1 with CL = 250ms;

carelease_case2a = A20(:,6); %Recordings of JSR released calcium ion concentration of each heart beat in case 2 with CL = 250ms;

carelease_case3a = A20(:,7); %Recordings of JSR released calcium ion concentration of each heart beat in case 3 with CL = 250ms;

carelease_cona = A20(:,8); %Recordings of JSR released calcium ion concentration of each heart beat in control condition with CL = 250ms;

casr_case1b = A21(:,1); %Recordings of SR calcium ion concentration of each heart beat in case 1 with CL = 300ms;

casr_case2b = A21(:,2); %Recordings of SR calcium ion concentration of each heart beat in case 2 with CL = 300ms;

casr_case3b = A21(:,3); %Recordings of SR calcium ion concentration of each heart beat in case 3 with CL = 300ms;

casr_conb = A21(:,4); %Recordings of SR calcium ion concentration of each heart beat in control condition with CL = 300ms;

carelease_case1b = A21(:,5); %Recordings of JSR released calcium ion concentration of each heart beat in case 1 with CL = 300ms;

carelease_case2b = A21(:,6); %Recordings of JSR released calcium ion concentration of each heart beat in case 2 with CL = 300ms;

carelease_case3b = A21(:,7); %Recordings of JSR released calcium ion concentration of each heart beat in case 3 with CL = 300ms;

carelease_conb = A21(:,8); %Recordings of JSR released calcium ion concentration of

each hear beat in control condition with CL = 300ms;

```
%cell geometry
```

```
L=0.01;
```

```
rad=0.0011;
```

```
vcell=1000*3.14*rad*rad*L;
```

```
Ageo=2*3.14*rad*rad+2*3.14*rad*L;
```

```
Acap=2*Ageo;
```

```
vmyo=0.68*vcell;
```

```
vnsr=0.0552*vcell;
```

```
vjsr=0.0048*vcell;
```

```
r_v = vjsr/(vnsr+vjsr);
```

```
plot(casr_case1a(91:638),carelease_case1a(91:638)./casr_case1a(91:638)*r_v,'r','linewidth',1);
```

```
hold on
```

```
plot(casr_case1a(639:1000),carelease_case1a(639:1000)./casr_case1a(639:1000)*r_v,'r.');
```

```
hold on
```

```
plot(casr_case2a(35:277),carelease_case2a(35:277)./casr_case2a(35:277)*r_v,'b','linewidth',1);
```

```
hold on
```

```
plot(casr_case2a(278:1000),carelease_case2a(278:1000)./casr_case2a(278:1000)*r_v,'bo');
```

```
hold on
```

```
plot(casr_case3a(19:166),carelease_case3a(19:166)./casr_case3a(19:166)*r_v,'m','linewidth',1);
```

```
hold on
```

```
plot(casr_case3a(167:1000),carelease_case3a(167:1000)./casr_case3a(167:1000)*r_v,'mx');
```

```
hold on
```

```
plot(casr_cona(8:128),carelease_cona(8:128)./casr_cona(8:128)*r_v,'g','linewidth',1);
```

```
xlabel('{SR Ca2+ content (mmol/L)}');
```

```
ylabel('{fractional SR Ca2+ release (%)}');
```

```
legend2 = legend('case 1','case 1','case 2','case 2','case 3','case 3','control');
```

```
set(legend2,'Position',[0.154 0.705 0.173 0.184]);
```

```
%axes('position',[0.2 0.6 0.15 0.3]);
```

```
axes('position',[0.79 0.62 0.08 0.12],'YTick',[0.106 0.112],'XTick',[2.7 2.8])
```

```
plot(casr_case1a(277:535),carelease_case1a(277:535)./casr_case1a(277:535)*r_v,'r','linewidth',1);
```

```
hold on
```

```
plot(casr_case2a(102:185),carelease_case2a(102:185)./casr_case2a(102:185)*r_v,'b','linewidth'
```

```

,1);
hold on
plot(casr_case3a(54:90),carelease_case3a(54:90)./casr_case3a(54:90)*r_v,'m','linewidth',1);
hold on
plot(casr_cona(25:40),carelease_cona(25:40)./casr_cona(25:40)*r_v,'g','linewidth',1);
axis([2.7 2.8 0.106 0.112]);

annotation('textbox',[0.71 0.87 0.174 0.044],'String',{'CL = 250ms'},'FitBoxToText','off');

subplot(2,1,2)
plot(casr_case1b(97:1000),carelease_case1b(97:1000)./casr_case1b(97:1000)*r_v,'r','linewidth'
,1);
hold on
plot(casr_case2b(27:267),carelease_case2b(27:267)./casr_case2b(27:267)*r_v,'b','linewidth',1);
hold on
plot(casr_case2b(268:1000),carelease_case2b(268:1000)./casr_case2b(268:1000)*r_v,'b. ');
hold on
plot(casr_case3b(14:156),carelease_case3b(14:156)./casr_case3b(14:156)*r_v,'m','linewidth',1);
hold on
plot(casr_case3b(157:1000),carelease_case3b(157:1000)./casr_case3b(157:1000)*r_v,'mx');
hold on
plot(casr_conb(5:95),carelease_conb(5:95)./casr_conb(5:95)*r_v,'g','linewidth',1);

xlabel('{SR Ca^2^+ content (mmol/L)}');
ylabel('{fractional SR Ca^2^+ release (%)}');

legend1 = legend('case 1','case 2','case 2','case 3','case 3','control');
set(legend1,'Position',[0.154 0.226 0.173 0.213]);
axes('position',[0.79 0.15 0.08 0.12],'YTick',[0.126 0.132],'XTick',[2.7 2.8]);
plot(casr_case1b(513:1000),carelease_case1b(513:1000)./casr_case1b(513:1000)*r_v,'r','linewi
dth',1);
hold on
plot(casr_case2b(137:268),carelease_case2b(137:268)./casr_case2b(137:268)*r_v,'b','linewidth
',1);
hold on
plot(casr_case3b(59:100),carelease_case3b(59:100)./casr_case3b(59:100),'m','linewidth',1);
hold on
plot(casr_conb(25:37),carelease_conb(25:37)./casr_conb(25:37)*r_v,'g','linewidth',1);

axis([2.7 2.8 0.126 0.132]);
annotation('textbox',[0.718 0.395 0.173 0.044],'String',{'CL = 300ms'},'FitBoxToText','off');
annotation('textbox',[0.023 0.93 0.059 0.044],'String',{'a'},'EdgeColor',[1 1 1]);
annotation('textbox',[0.023 0.465 0.059 0.044],'String',{'b'},'EdgeColor',[1 1 1]);

```

%figure3.m

%%%

%%%figure3.m %%

%%%

clear all;

close all;

clc;

subplot(2,1,1)

load A20; %Its contents are explained in the following

load A21; %Its contents are explained in the following

casr_case1a = A20(:,1); %Recordings of SR calcium ion concentration of each heart beat(the total number is 1000) in case 1 with CL = 250ms;

casr_case2a = A20(:,2); %Recordings of SR calcium ion concentration of each heart beat in case 2 with CL = 250ms;

casr_case3a = A20(:,3); %Recordings of SR calcium ion concentration of each heart beat in case 3 with CL = 250ms;

casr_cona = A20(:,4); %Recordings of SR calcium ion concentration of each heart beat in control condition with CL = 250ms;

carelease_case1a = A20(:,5); %Recordings of JSR released calcium ion concentration of each heart beat in case 1 with CL = 250ms;

carelease_case2a = A20(:,6); %Recordings of JSR released calcium ion concentration of each heart beat in case 2 with CL = 250ms;

carelease_case3a = A20(:,7); %Recordings of JSR released calcium ion concentration of each heart beat in case 3 with CL = 250ms;

carelease_cona = A20(:,8); %Recordings of JSR released calcium ion concentration of each heart beat in control condition with CL = 250ms;

casr_case1b = A21(:,1); %Recordings of SR calcium ion concentration of each heart beat in case 1 with CL = 300ms;

casr_case2b = A21(:,2); %Recordings of SR calcium ion concentration of each heart beat in case 2 with CL = 300ms;

casr_case3b = A21(:,3); %Recordings of SR calcium ion concentration of each heart beat in case 3 with CL = 300ms;

casr_conb = A21(:,4); %Recordings of SR calcium ion concentration of each heart beat in control condition with CL = 300ms;

carelease_case1b = A21(:,5); %Recordings of JSR released calcium ion concentration of each heart beat in case 1 with CL = 300ms;

carelease_case2b = A21(:,6); %Recordings of JSR released calcium ion concentration of each heart beat in case 2 with CL = 300ms;

carelease_case3b = A21(:,7); %Recordings of JSR released calcium ion concentration of each heart beat in case 3 with CL = 300ms;

carelease_conb = A21(:,8); %Recordings of JSR released calcium ion concentration of

each hear beat in control condition with CL = 300ms;

```
%cell geometry
```

```
L=0.01;
```

```
rad=0.0011;
```

```
vcell=1000*3.14*rad*rad*L;
```

```
Ageo=2*3.14*rad*rad+2*3.14*rad*L;
```

```
Acap=2*Ageo;
```

```
vmyo=0.68*vcell;
```

```
vnsr=0.0552*vcell;
```

```
vjsr=0.0048*vcell;
```

```
r_v = vjsr/(vnsr+vjsr);
```

```
plot(casr_case1a(91:638),carelease_case1a(91:638)./casr_case1a(91:638)*r_v,'r','linewidth',1);
```

```
hold on
```

```
plot(casr_case1a(639:1000),carelease_case1a(639:1000)./casr_case1a(639:1000)*r_v,'r');
```

```
hold on
```

```
plot(casr_case2a(35:277),carelease_case2a(35:277)./casr_case2a(35:277)*r_v,'b','linewidth',1);
```

```
hold on
```

```
plot(casr_case2a(278:1000),carelease_case2a(278:1000)./casr_case2a(278:1000)*r_v,'bo');
```

```
hold on
```

```
plot(casr_case3a(19:166),carelease_case3a(19:166)./casr_case3a(19:166)*r_v,'m','linewidth',1);
```

```
hold on
```

```
plot(casr_case3a(167:1000),carelease_case3a(167:1000)./casr_case3a(167:1000)*r_v,'mx');
```

```
hold on
```

```
plot(casr_cona(8:128),carelease_cona(8:128)./casr_cona(8:128)*r_v,'g','linewidth',1);
```

```
xlabel('{SR Ca2+ content (mmol/L)}');
```

```
ylabel('{fractional SR Ca2+ release (%)}');
```

```
legend2 = legend('case 1','case 1','case 2','case 2','case 3','case 3','control');
```

```
set(legend2,'Position',[0.154 0.705 0.173 0.184]);
```

```
%axes('position',[0.2 0.6 0.15 0.3]);
```

```
axes('position',[0.79 0.62 0.08 0.12],'YTick',[0.106 0.112],'XTick',[2.7 2.8])
```

```
plot(casr_case1a(277:535),carelease_case1a(277:535)./casr_case1a(277:535)*r_v,'r','linewidth',1);
```

```
hold on
```

```
plot(casr_case2a(102:185),carelease_case2a(102:185)./casr_case2a(102:185)*r_v,'b','linewidth'
```

```

,1);
hold on
plot(casr_case3a(54:90),carelease_case3a(54:90)./casr_case3a(54:90)*r_v,'m','linewidth',1);
hold on
plot(casr_cona(25:40),carelease_cona(25:40)./casr_cona(25:40)*r_v,'g','linewidth',1);
axis([2.7 2.8 0.106 0.112]);

annotation('textbox',[0.71 0.87 0.174 0.044],'String',{'CL = 250ms'},'FitBoxToText','off');

subplot(2,1,2)
plot(casr_case1b(97:1000),carelease_case1b(97:1000)./casr_case1b(97:1000)*r_v,'r','linewidth'
,1);
hold on
plot(casr_case2b(27:267),carelease_case2b(27:267)./casr_case2b(27:267)*r_v,'b','linewidth',1);
hold on
plot(casr_case2b(268:1000),carelease_case2b(268:1000)./casr_case2b(268:1000)*r_v,'b. ');
hold on
plot(casr_case3b(14:156),carelease_case3b(14:156)./casr_case3b(14:156)*r_v,'m','linewidth',1);
hold on
plot(casr_case3b(157:1000),carelease_case3b(157:1000)./casr_case3b(157:1000)*r_v,'mx');
hold on
plot(casr_conb(5:95),carelease_conb(5:95)./casr_conb(5:95)*r_v,'g','linewidth',1);

xlabel('{SR Ca^2^+ content (mmol/L)}');
ylabel('{fractional SR Ca^2^+ release (%)}');

legend1 = legend('case 1','case 2','case 2','case 3','case 3','control');
set(legend1,'Position',[0.154 0.226 0.173 0.213]);
axes('position',[0.79 0.15 0.08 0.12],'YTick',[0.126 0.132],'XTick',[2.7 2.8]);
plot(casr_case1b(513:1000),carelease_case1b(513:1000)./casr_case1b(513:1000)*r_v,'r','linewi
dth',1);
hold on
plot(casr_case2b(137:268),carelease_case2b(137:268)./casr_case2b(137:268)*r_v,'b','linewidth
',1);
hold on
plot(casr_case3b(59:100),carelease_case3b(59:100)./casr_case3b(59:100),'m','linewidth',1);
hold on
plot(casr_conb(25:37),carelease_conb(25:37)./casr_conb(25:37)*r_v,'g','linewidth',1);

axis([2.7 2.8 0.126 0.132]);
annotation('textbox',[0.718 0.395 0.173 0.044],'String',{'CL = 300ms'},'FitBoxToText','off');
annotation('textbox',[0.023 0.93 0.059 0.044],'String',{'a'},'EdgeColor',[1 1 1]);
annotation('textbox',[0.023 0.465 0.059 0.044],'String',{'b'},'EdgeColor',[1 1 1]);

```

%figure4.m

%%%

%%%

%%%

clear all;

close all;

clc;

load A4; % Recordings of the values of Cansr, Cai and their ratios when alternans begin in different ischemic cases.

cansr = A4(:,1);

ph65 = 'Case 1';

ph66 = 'Case 2';

ph67 = 'Case 3';

case1a = repmat(ph65,4,1); %char. axis label

case2a = repmat(ph66,13,1); %char. axis label

case3a = repmat(ph67,8,1); %char. axis label

G = char(case1a,case2a,case3a);

subplot(2,2,1);

m=boxplot(cansr,G);

set(m,'LineWidth',1);

xlabel('Ischemic conditions');

ylabel('Ca_{n_s_r} (mmol/L)');

cai = A4(:,2);

ph65 = 'Case 1';

ph66 = 'Case 2';

ph67 = 'Case 3';

case1b = repmat(ph65,4,1);

case2b = repmat(ph66,13,1);

case3b = repmat(ph67,8,1);

G = char(case1b,case2b,case3b);

subplot(2,2,2);

m=boxplot(cai,G);

set(m,'LineWidth',1);

xlabel('Ischemic conditions');

ylabel('Ca_i (mmol/L)');

ratio = A4(:,3);

ph65 = 'Case 1';

ph66 = 'Case 2';

```
ph67 = 'Case 3';  
case1c = repmat(ph65,4,1);  
case2c = repmat(ph66,13,1);  
case3c = repmat(ph67,8,1);  
G = char(case1c,case2c,case3c);  
subplot(2,2,3);  
m=boxplot(ratio,G);  
set(m,'LineWidth',1);  
xlabel('Ischemic conditions');  
ylabel('Ca_n_s_r/Ca_i');
```



```
%figure5.m
```

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
```

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
```

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
```

```
clear all;
```

```
close all;
```

```
clc;
```

```
load A5; %There are instructions about its content in the following.
```

```
casrN = A5(:,1)./max(A5(:,1)); %normalized SR calcium ion concentration
```

```
caiN = A5(:,2)./max(A5(:,2)); %normalized cytoplasmic calcium ion concentration
```

```
ratio = A5(:,3)/A5(:,2); %A(:,3) represents the NSR calcium ion concentration. This value  
shows the ratio of calcium ion concentration between NSR to the cytoplasm.
```

```
subplot(2,1,1)
```

```
plot(casrN,',' , 'Markersize',4);
```

```
hold on
```

```
plot(caiN,',' , 'Color',[0.85 0.16 0], 'Markersize',4, 'LineWidth',1);
```

```
xlabel('Number of Beats');
```

```
set(subplot(2,1,1), 'LineWidth',1);
```

```
legend1 = legend('Normalized diastolic Ca_s_r', 'Normalized diastolic Ca_i');
```

```
set(legend1, 'Position',[0.182 0.6 0.368 0.1], 'LineWidth',1);
```

```
B1=300:330;
```

```
axes('position',[0.76 0.65 0.11 0.155]);
```

```
plot(B1,casrN(300:330),',' );
```

```
axis([300 330 0.82 0.84]);
```

```
set(axes, 'LineWidth',1);
```

```
annotation('textbox',[0.145 0.85 0.162 0.0475], 'String',{'CL =  
250ms'}, 'FitBoxToText','off', 'LineWidth',1);
```

```
subplot(2,1,2)
```

```
plot(ratio,',' , 'Markersize',4, 'LineWidth',1);
```

```
xlabel('Number of Beats');
```

```
ylabel('Ca_n_s_r/Ca_i');
```

```
annotation('textbox',[0.145 0.37 0.162 0.0475], 'String',{'CL =  
250ms'}, 'FitBoxToText','off', 'LineWidth',1);
```

```
set(subplot(2,1,2), 'LineWidth',1);
```

```
annotation('textbox',[0.029 0.9 0.059 0.067], 'String',{'a'}, 'LineStyle','none');
```

```
annotation('textbox',[0.029 0.466 0.059 0.067], 'String',{'b'}, 'LineStyle','none');
```