

TR_1D_model1_SS\TR_1D_model1_func_calc_b_int

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```

% TR_1D_model1_SS\TR_1D_model1_func_calc_b_int.m
%
% function [b_int,bJac_int,iflag] = ...
%   TR_1D_model1_calc_b_int(x_state,epsilon,Param);
%
% This function calculates the b vector and its
% Jacobian bJac of a DAE system of the form :
%
%   epsilon(k) * df_dt(k) = b(k) -
%       sum_{j} {A(k,j)*x_state(j)}
%
% for a reaction system that is comprised of an
% arbitrary number of species and an arbitrary number
% of reactions. The vector epsilon is used as an
% integer mask to tell the solver where to calculate
% the elements of b and bJac from the reaction model
% for the interior points. The reaction system is
% comprised of an arbitrary number of reactions with
% specified stoichiometric coefficients and exponential
% rate law powers for each species. This procedure
% goes through each interior grid point and passes the
% concentration and temperature values at this point
% to a general subroutine that evaluates the source
% term and its derivatives at this point. Then, a
% shifting routine takes this data and places it in the
% appropriate spots in the b vector and bJac matrix.
% Since the reaction rate depends on the values of the
% concentration and temperature fields at a single
% point, this routine may be used in any geometry.
%
% INPUT :
% =====
% x_state      REAL(num_DOF) where
%              num_DOF = Grid.num_pts *
%              (ProbDim.num_species + 1)
%              This is the column vector of the state
%              variables containing the concentration
%              and temperature profiles.
% epsilon      INT(num_DOF)
%              This is a vector of integers that has
%              1's at the interior point equations
%              and 0's at the boundary point
%              equations.
% Param        This data structure contains the
%              system parameters stored in the
%              fields :
%              .ProbDim

```

```

%           .Reactor
%           .Rxn
%           .Physical
%           .Grid
%           see TR_1D_model1_SS.m for details of
%           these data structures
%
% OUTPUT :
% =====
% b_int      REAL(num_DOF)
%           This column vector contains the source
%           terms at the interior points of the
%           concentration and temperature PDE's
% bJac_int   REAL(num_DOF,num_DOF)
%           This is the Jacobian element of b_int
%           whose (m,n) element is defined as the
%           partial derivative of b_int(m) with
%           respect to x_state(n).
%
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% Massachusetts Institute of Technology
% Department of Chemical Engineering
% 7/2/2001
%
% Version as of 7/25/2001

```

```

function [b_int,bJac_int,iflag] = ...
    TR_1D_model1_calc_b_int(x_state,epsilon,Param);

```

```

iflag = 0;

```

```

func_name = 'TR_1D_model1_calc_b_int';

```

```

% This sets what to do in case an assertion fails.

```

```

i_error = 2;

```

```

% First, check input.

```

```

% First, extract parameter structures.

```

```

if(~isstruct(Param))
    iflag = -1;
    message = [func_name, ': ', ...
              'Paramis not a structure'];
    if(i_error ~= 0)
        if(i_error > 1)
            save dump_error.mat;
        end

```

```
    error(message);
else
    b_int = 0; bJac_int=0;
    return;
end
end

% ProbDim
if(~isfield(Param,'ProbDim'))
    iflag = -1;
    message = [func_name, ': ', ...
        'Param does not contain ProbDim'];
    if(i_error ~= 0)
        if(i_error > 1)
            save dump_error.mat;
        end
        error(message);
    else
        b_int=0; bJac_int=0;
        return;
    end
end
ProbDim = Param.ProbDim;

% Reactor
if(~isfield(Param,'Reactor'))
    iflag = -1;
    message = [func_name, ': ', ...
        'Param does not contain Reactor'];
    if(i_error ~= 0)
        if(i_error > 1)
            save dump_error.mat;
        end
        error(message);
    else
        b_int=0; bJac_int=0;
        return;
    end
end
Reactor = Param.Reactor;

% Rxn
if(~isfield(Param,'Rxn'))
    iflag = -1;
    message = [func_name, ': ', ...
        'Param does not contain Rxn'];
    if(i_error ~= 0)
        if(i_error > 1)
            save dump_error.mat;
        end
    end
end
```

```
        error(message);
    else
        return;
    end
end
Rxn = Param.Rxn;

% Physical
if(~isfield(Param,'Physical'))
    iflag = -1;
    message = [func_name, ': ', ...
              'Param does not contain Physical'];
    if(i_error ~= 0)
        if(i_error > 1)
            save dump_error.mat;
        end
        error(message);
    else
        return;
    end
end
Physical = Param.Physical;

% Grid
if(~isfield(Param,'Grid'))
    iflag = -1;
    message = [func_name, ': ', ...
              'Param does not contain Grid'];
    if(i_error ~= 0)
        if(i_error > 1)
            save dump_error.mat;
        end
        error(message);
    else
        return;
    end
end
Grid = Param.Grid;

% Extract the proper dimension of x_state and epsilon.

num_DOF = (ProbDim.num_species+1)*Grid.num_pts;

% check x_state and epsilon to be vectors of proper type

% x_state
dim = num_DOF; check_column = 1;
check_real = 1; check_sign = 0; check_int = 0;
```

```
assert_vector(i_error,x_state,'x_state',func_name, ...
  dim,check_real,check_sign,check_int,check_column);
```

```
% epsilon
```

```
dim = num_DOF; check_column = 1;
check_real = 1; check_sign = 2; check_int = 1;
assert_vector(i_error,epsilon,'epsilon', func_name, ...
  dim,check_real,check_sign,check_int,check_column);
```

```
%PDL> Initialize b, bJac to zeros
```

```
b_int = linspace(0,0,num_DOF)';
```

```
max_nonzero = (ProbDim.num_species+1)*Grid.num_pts;
bJac_int = spalloc(num_DOF,num_DOF,max_nonzero);
```

```
%PROCEDURE: unstack_state
```

```
%PDL> Unstack x_state to state_data variable format
```

```
%ENDPROCEDURE
```

```
[State,iflag_func] = unstack_state(x_state, ...
  ProbDim.num_species,Grid.num_pts);
```

```
if(iflag_func <= 0)
```

```
  iflag = -2;
```

```
  message = [func_name, ': ', ...
    'Error (', int2str(iflag_func), ') ', ...
    'returned from unstack_state'];
```

```
  if(i_error ~= 0)
```

```
    if(i_error > 1)
```

```
      save dump_error.mat;
```

```
    end
```

```
    error(message);
```

```
  else
```

```
    return;
```

```
  end
```

```
end
```

```
%PDL> FOR EVERY interior grid point ipoint
```

```
%   FROM 1 TO Grid.num_pts
```

```
%   WHERE epsilon(ipoint) is non-zero
```

```
for ipoint = 1:Grid.num_pts
```

```
if(epsilon(ipoint) ~= 0)
```

```
%   PROCEDURE: reaction_network_model
```

```
%   PDL> Pass the concentrations and temperature
```

```
%   at this point along with the density and
```

```

%   heat_capacity to a general routine that
%   implements the general reaction network model
%   and returns the reaction rates, the partial
%   derivatives of the rates with respect to each
%   concentration and temperature.
%   ENDPROCEDURE

conc_loc = State.conc(ipoint,:);
Temp_loc = State.Temp(ipoint);

% the local reaction is characterized by the data
% in a structure RxnRate that contains the following
% fields :
% .rxn_time_deriv_c - the total time derivative of each
%                   concentration due to all reactions
% .rxn_time_deriv_T - the total time derivative of the
%                   temperature due to all reactions
% .rxn_rate - the rates of each reaction
% .rxn_rate_deriv_c - the derivative of each reaction rate
%                   with respect to each concentration
% .rxn_rate_deriv_T - the derivative of each reaction rate
%                   with respect to the temperature

[RxnRate_loc, iflag_func] = ...
    reaction\_network\_model(...
    ProbDim.num_species,ProbDim.num_rxn, ...
    conc_loc,Temp_loc,Rxn,Physical.density,Physical.Cp);
if(iflag_func <= 0)
    iflag = -3;
    message = [func_name, ': ', ...
              'Error (', int2str(iflag_func), ') ', ...
              'returned from reaction_network_model'];
    if(i_error ~= 0)
        if(i_error > 1)
            save dump_error.mat;
        end
        error(message);
    else
        return;
    end
end

%   PROCEDURE: shift_rxn_source_term
%   PDL> Next, b and bJac are incremented by a
%   procedure that takes as input the grid point
%   and the reaction rates and their derivatives,
%   and adds these results to the appropriate b
%   and bJac locations.
%   ENDPROCEDURE

```

```
[b_loc,bJac_loc,iflag_func] = shift\_rxn\_source\_term(...  
    ProbDim,Grid,Rxn,Physical,ipoint,RxnRate_loc);  
if(iflag_func <= 0)  
    iflag = -4;  
    message = [func_name, ': ', ...  
              'Error (', int2str(iflag_func), ') ', ...  
              'returned from shift_rxn_source_term'];  
    if(i_error ~= 0)  
        if(i_error > 1)  
            save dump_error.mat;  
        end  
        error(message);  
    else  
        return;  
    end  
end  
end
```

```
b_int = b_int + b_loc;  
bJac_int = bJac_int + bJac_loc;
```

%PDL> ENDFOR for loop over interior points

```
end  
end
```

%PDL> Add contribution from heat transfer to the
% jacket

```
[b_HT,bJac_HT,iflag_func] = jacket\_heat\_transfer( ...  
    State,epsilon,ProbDim,Reactor,Physical,Grid);  
if(iflag_func <= 0)  
    iflag = -5;  
    message = [func_name, ': ', ...  
              'Error (', int2str(iflag_func), ') ', ...  
              'returned from jacket_heat_transfer'];  
    if(i_error ~= 0)  
        if(i_error > 1)  
            save dump_error.mat;  
        end  
        error(message);  
    else  
        return;  
    end  
end  
end
```

```
b_int = b_int + b_HT;  
bJac_int = bJac_int + bJac_HT;
```

iflag = 1;

return;