

About mfiles for HydroBase3:

The basic read function is `[profiles, nsta] = hb_read_file(dir, root, extent);`

INPUT: 1 or 3 strings that are concatenated to specify a filename.

OUTPUT: **profiles** is a struct array with the fields listed below

nsta : length(profiles)

Returns a structure with the following form:

```
% profile.country
%   .ship,
%   .cruise
%   .station
%   .year
%   .month
%   .day
%   .orig
%   .instr
%   .lat
%   .lon
%   .bdpth
%   .nobs
%   .nprops
%   .ms10
%   .ms1
%   .prop_id ( names of variable fields in profile.data)
%   .qual
%   .data (structure containing nprops separate fields )
%   .pr
%   .te (etc....)
```

profiles(1).station stores the station #

profiles(1).nobs stores number of observation levels in the profile

profiles(1).nprops stores number of properties

profiles(1).prop_id lists the names of available properties

property values are stored as column vectors in the field data which is a struct array:

```
profiles(1).data.pr    % pressure
profiles(1).data.t90   % in situ temperature
profiles(1).data.sa    % salinity
profiles(1).data.o2    % oxygen ( in micromoles/kg)
*****
```

The basic write function is :

```
function err = hb_write_file(profiles, fname, mode)
% Creates file for writing in specified mode, writes profiles
% and closes the file.
% profiles is struct array of profiles to be written
% fname is full pathname of file
% mode is 0 (zero) or 'w' for Overwrite
%       1 NoClobber
%       2 or 'a' for Append
% Returns err > 0 for an error.
```

which performs the following:

```
fid = hb_create_file(fname, 0);
[m,nsta] = size(profiles);
for ii=1:nsta
```

```

    err = hb_write_profile(fid,profiles(ii));
end
fclose(fid);

```

Useful ways of searching through the data.....

```

[P,nsta]=hb_read_file('2008may.hb2_ctd');

```

```

staid = [P(:).station]; % returns array of all station #s
lats = [P(:).lat];      % returns array of all station latitudes

```

```

indx = find(staid==9009); % search for a particular station
OR
indx = find([P(:).station]==9014)

```

% list properties available for a particular profile:

P(indx).prop_id

Returns names (strings) of each property:

```

pr
de
t90
sa
o2

```

These strings form the fields of the structure **data** which is itself a field of **P()**

P(indx).data.pr -- array of pressure for station # 9014

P(indx).data.de -- array of depths

propname = P(indx).prop_id(3) -- name of 3rd property in station #9014

P(indx).data.(propname) -- (**propname**) denotes a dynamic variable : propname is a string containing name of variable.

%Load properties from a single station

```

indx= find( [P(:).station] == 9009);

```

```

pr = P(indx).data.pr;

```

```

t90 = P(indx).data.t90;

```

```

sa = P(indx).data.sa);

```

% Compute potential temperature

```

th= sw_ptmp(sa, t90, pr, 0.0);

```

% Load each property from all profiles into a matrix

```

pr=P(1).data.pr;

```

```

t90=P(1).data.t90;

```

```

sa=P(1).data.sa;

```

```

o2=P(1).data.o2;

```

```

for ii=2:nsta

```

```

    pr=merge(pr,P(ii).data.pr);

```

```

    t90=merge(te,P(ii).data.t90);

```

```

    sa=merge(sa,P(ii).data.sa);

```

```

    if ~ isempty(strmatch('o2 ',P(ii).prop_id) % checks if the profile has o2 data

```

```

        o2=merge(o2,P(ii).data.o2);

```

```

    else

```

```

        o2=merge(o2,[NaN]);

```

```

    end

```

```

end

```