

# Input format of FRENDY

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# Input format of FRENDY

- FRENDY treats two types of the input formats.
  - FRENDY original input format
  - NJOY compatible
- Simple input format
  - Nuclear data file name and processing mode are **only** required for the processing.
    - FRENDY has recommended parameters in the source code.
    - User can also change (override) parameters.

# Input format of FRENDY and NJOY

- Input parameters of FRENDY consist of “input data name” and “input data”.
- Comment line is similar to C/C++.
- //~ or /\* ~ \*/
- Input format of FRENDY is easy to understand.
- This input format is suitable for beginners.

## [Sample input of FRENDY]

```
ace_fast_mode // Processing mode
nucl_file_name U235.dat
ace_file_name U235.ace
temp 296.0
```

## [Sample input of NJOY]

```
reconr / command
20 21 / input(tape20), output(tape21)
'pendf tape for JENDL-4 U235' / identifier for PENDF
9228 / mat
1.00e-03 0.00 / err, temp
0 /
broadr / command
20 21 22 / endf, pendf(in), pendf(out)
9228 1 / mat, temp no
1.00e-03 -5.0E+2 / err, thnmax
296.0 / temp
0 /
gaspr / command
20 22 23 / endf, pendf(in), pendf(out)
purr / command
20 23 25 / endf, pendf(in), pendf(out)
9228 1 7 20 500 / mat, temp no, sig no, bin no, lad no
296.0 / temp
1E10 1E4 1E3 300 100 30 10 / sig zero
0 /
acer / command
20 25 0 30 31 / nendf, npend, ngend, nace, ndir
1 1 1 0.30 / iopt(fast), iprint(max), itype, suffix
'ACE file for JENDL-4 U235' / descriptive character
9228 296.0 / mat, temp
1 1 / newfor(yes), iopp(yes)
1 1 1 / thin(1), thin(2), thin(3)
stop /
```

# FRENDY original input format

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# Basics of FRENDY input format

- First line **must be** processing mode.
  - ace\_fast\_mode : Neutron induced ACE file
  - ace\_tsl\_mode : ACE file of thermal scattering law data
  - ace\_dosi\_mode: Dosimetry ACE file
- Other lines are free format.
  - Setting “input data name” and “input data”
  - Bracket is used to describe array data, e.g., (1.0 2.0 3.0).
  - Text data is surrounded by single or double quotation mark, e.g., “~” or ‘~’.
    - Multiline is available for array data and text data.
- Comment line is similar to C/C++.
  - //~ or /\* ~ \*/

# Main input parameters

- Required parameters
  - nucl\_file\_name : Evaluated nuclear data file name
  - nucl\_file\_name\_tsl : TSL data file name (TSL only)
- Other main input parameters
  - temp : Temperature (K) (Default: 293.6 K)
  - ace\_file\_name : ACE file name
  - ace\_dir\_file\_name : XSDIR file name
  - suffix\_id : Suffix ID of ACE file
  - ace\_label\_data : Comment line of ACE file (one line)
  - thermal\_z\_a\_id\_name :  $S(\alpha, \beta)$  identifier of MCNP (TSL only)
- Additional parameters
  - Write\_pendf\_probability\_table : PENDF file name after probability table generation
  - Write\_pendf\_tsl : PENDF file name after TSL data processing (TSL only)
    - PENDF (Point-wise ENDF) file is a processing result of NJOY.
    - These PENDF file can be used as input PENDF of NJOY.

# Example of FRENDY input (Neutron induced)

```
ace_fast_mode // Processing mode
nucl_file_name    ../lib/U235.dat
temp /* [K] */    300.0
ace_file_name     ./ace/U235.ace
ace_dir_file_name ./xsd/U235.xsdir
ace_label_data    "U-235 from JENDL-4.0"
suffix_id         0.50
```

First line **must be** set  
processing mode.

Relative path or  
absolute path

- Processing conditions of above example
  - Evaluated nuclear data file name:   ../lib/U235.dat
  - Temperature:                       300.0 [K]
  - ACE file name:                     ./ace/U235.ace
  - XSDIR file name:                  ./xsd/U235.xsdir
  - Comment line of ACE file:        U-235 from JENDL-4.0
  - Suffix ID:                         0.50

# Modification of processing conditions (Neutron induced)

- Please try to modify input file if processing conditions are changed as follows:
  - Evaluated nuclear data file name: `./j40/lib/Fe056.dat`
  - Temperature: `550.0 [K]`
  - ACE file name: `./j40/ace/Fe056.ace`
  - XSDIR file name: `./j40/xsd/Fe056.xsdir`
  - Comment line of ACE file: `Fe-056 from JENDL-4.0`
  - Suffix ID: `0.10`
  - PENDF file name: `./j40/pendf/Fe056.pendf`

The answer is next slide.



# Modification of processing conditions (Neutron induced)

Input is modified to meet the processing conditions.

```
ace_fast_mode // Processing mode
nucl_file_name ./j40/lib/Fe056.dat
temp /* [K] */ 550.0
ace_file_name ./j40/ace/Fe056.ace
ace_dir_file_name ./j40/xsd/Fe056.xsdir
ace_label_data 'Fe-056 from JENDL-4.0'
suffix_id 0.10
write PENDF probability_table ./j40/PENDF/Fe056.PENDF
```

Text data is surrounded by single or double quotation mark.

Adding PENDF output option

# Example of FRENDY input (TSL)

```
ace_tsl_mode // Processing mode
nucl_file_name      ../lib/H001.dat
nucl_file_name_tsl  ../lib_sab/01_h_in_h2o.txt
ace_label_data      "HinH2O from JENDL-4.0"
temp                296.0
ace_file_name        ./ace_sab/lwtr.ace
ace_dir_file_name    ./xsd_sab/lwtr.xsdir
suffix_id            0.50
thermal_zs_id_name   "lwtr"
```

First line **must be** set processing mode.

Temperature prepared in TSL data file (nucl\_file\_name\_tsl) is **only** available.

S( $\alpha$ ,  $\beta$ ) identifier of MCNP (Maximum: 6 characters)

- Processing conditions of above example
  - TSL data file name: ../lib\_sab/01\_h\_in\_h2o.txt
  - S( $\alpha$ ,  $\beta$ ) identifier of MCNP: lwtr
    - lwtr: light water
    - This name is used in S( $\alpha$ ,  $\beta$ ) card of MCNP input.

# Modification of processing conditions (TSL)

- Please try to modify input file if processing conditions are changed as follows:
  - Evaluated nuclear data file name: `./j40/lib/C000.dat`
  - TSL data file name: `./j40/lib/31_graphite.tx`
  - Temperature: `500.0 [K]`
  - ACE file name: `./j40/ace/graphite.ace`
  - XSDIR file name: `./j40/xsd/graphite.xsdir`
  - Comment line of ACE file: `Graphite from JENDL-4.0`
  - Suffix ID: `0.10`
  - S( $\alpha$ ,  $\beta$ ) identifier of MCNP: `grph`
  - PENDF file name: `./j40/pendf/graphite.pendf`

The answer is next slide.

# Modification of processing conditions (TSL)

Input is modified to meet the processing conditions.

```
ace_tsl_mode // Processing mode
nucl_file_name      ./j40/lib/C000.dat
nucl_file_name_tsl  ../lib_sab/31_graphite.txt
temp /* [K] */      500.0
ace_file_name       ./j40/ace/graphite.ace
ace_dir_file_name   ./j40/xsd/graphite.xsdir
ace_label_data      'Graphite from JENDL-4.0'
suffix_id           0.10
thermal_zaid_name   'grph'
write_pendf_tsl     ./j40/pendf/graphite.pendf
```

Adding PENDF output option

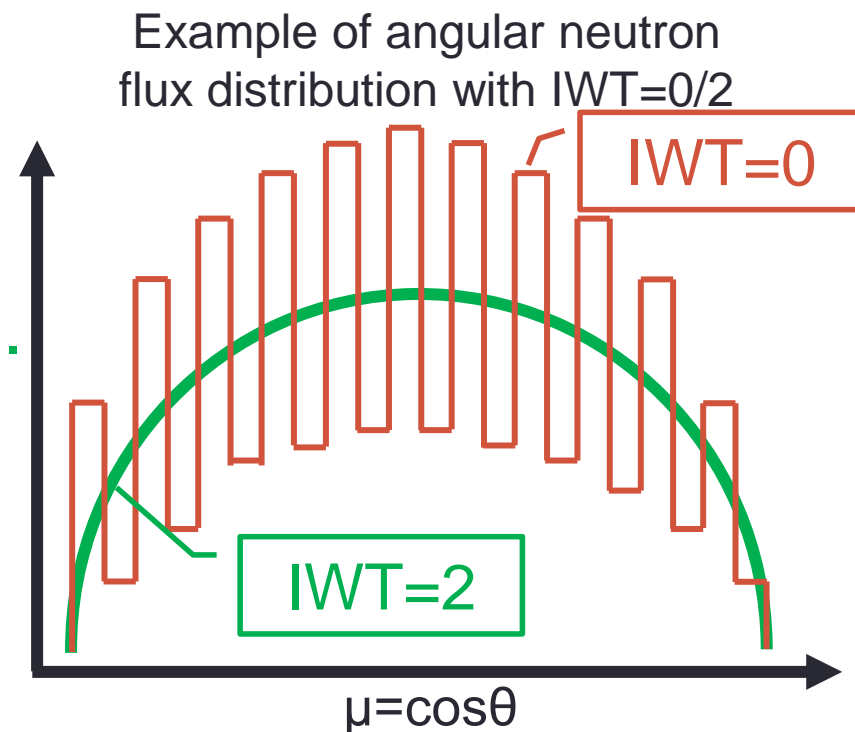
# TSL data generation for MCNP5 and PHITS 3.24

- Three types of ACE format is now available.
  - IFENG=0 (discrete), 1 (skewed), 2 (continuous)
- MCNP5 and PHITS ver.3.24 cannot treat IFENG=2.
  - The default input option of FRENDY version 2 generates IFENG=2.
  - Please add “weight\_option” and modify parameter from “tabulated” to “variable”.

```
ace_tsl_mode // Processing mode
nucl_file_name      ../lib/H001.dat
nucl_file_name_tsl  ../lib_sab/01_h_in_h2o.txt
ace_label_data      “HinH2O from JENDL-4.0”
temp               296.0
ace_file_name       ./ace_sab/lwtr.ace
ace_dir_file_name   ./xsd_sab/lwtr.xsdir
suffix_id           0.50
thermal_zs_id_name  “lwtr”
weight_option       variable
```

# Difference of IWT options

- IWT option affects the angular neutron flux distribution
  - Angular neutron flux distribution may become jagged.
- Recommend option: IWT=2
  - If users want to accurately calculate the angular neutron flux distribution.
  - IWT must be 2 for the multi-group cross section generation.
- Do not use IWT=1 (constant).
  - This option will affect not only the angular neutron flux distribution but also k-effective value.





# FRENDY original input format (Multi-group generation)

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# Input format of multi-group generation

- Input format of multi-group generation is similar to that of ACE file generation.
  - Difference is only **processing mode name**
  - Multi-group file can be generated from ENDF and ACE files
- Processing of U-235

```
mg_neutron_mode           //Process mode
nucl_file_name ( ../lib/U235.dat ) //ENDF file name
```

- Processing of H in H2O

```
mg_neutron_mode           //Process mode
nucl_file_name   ( ../lib/H001.dat ) //ENDF file name
nucl_file_name_tsl ( ../lib/01_h_in_h2o.txt )

mg_tsl_data_type  (hh2o) //This data type is used for MATXS
```

# Default option of multi-group generation

- Output format: **GENDF**
- Weighting spectrum: **1/E**
- Energy group structure: **XMAS 172-group structure**
- Flux calculation mode: **ultra-fine group slowing down calculation (120,000 groups)**
- Maximum Legendre order: **3 (P3)**

# Input Example (1/7): H-001

mg_neutron_mode		//Process mode
nucl_file_name	( ../lib/H001.dat )	//ENDF file name
mg_file_name	H001	//Output file name
mg_edit_option	( MATXS GENDF )	//Output format
temperature	300.0 //[K]	
mg_structure	( xmas_nea-lanl_172 )	//ign=18 in GROUPR/NJOY
mg_weighting_spectrum	( 1/e )	//iwt=3 in GROUPR/NJOY



Multi-group generation from ACE file

mg_neutron_mode		//Process mode
ace_file_name	( ../ace/H001.ace )	//ACE file name
mg_file_name	H001	//Output file name
mg_edit_option	( MATXS GENDF )	//Output format
mg_structure	( xmas_nea-lanl_172 )	//ign=18 in GROUPR/NJOY
mg_weighting_spectrum	( 1/e )	//iwt=3 in GROUPR/NJOY

# Input Example (2/7): U-235

```

mg_neutron_mode           //Process mode
nucl_file_name ( ../lib/U235.dat )    //ENDF file name
mg_file_name  U235              //Output file name
mg_edit_option ( SimpleGENDF  SimpleMATXS
                  GENDF  MATXS MGFlux )
//SimpleGENDF: MATXS format consistent with NJOY99
//SimpleMATXS: GENDF format consistent with NJOY99
//GENDF: GENDF format consistent with NJOY2016
//MATXS: MATXS format consistent with NJOY2016
temp           300.0 //[K]
mg_structure ( ign18 )           //ign=18 in GROUPR/NJOY
mg_weighting_spectrum ( iwt03 )  // iwt=3 in GROUPR/NJOY
legendre_order           3

```

# Input Example (3/7): U-235 with specified energy group structure and spectrum

```
mg_neutron_mode           //Process mode
nucl_file_name ( ../lib/U235.dat ) //ENDF file name
mg_file_name U235         //Output file name
mg_edit_option ( GENDF MATXS )
temp           300.0 //[K]
mg_structure ( 1.0E-5 0.625 5.0E5 1.0E6 2.0E+7 ) //4-groups
mg_weighting_spectrum ( 1.0E-5 1.0 2.0E+7 1.0 ) //constant
legendre_order           3
```

# Input Example (4/7): H in H<sub>2</sub>O

```
mg_neutron_mode           //Process mode
nucl_file_name      ( ../lib/H001.dat )      //ENDF file name
nucl_file_name_tsl  ( ../lib/01_h_in_h2o.txt ) //TSL file name
mg_edit_option      ( MATXS  GENDF )         //Output format
mg_file_name         H_in_H2O                 //Output file name
temperature          293.6 //[K]
mg_structure ( xmas_nea-lanl_172 )           //ign=18 in GROUPR/NJOY
mg_weighting_spectrum ( 1/e )                //iwt=3 in GROUPR/NJOY
mg_tsl_data_type     ( hh2o )
```



Multi-group generation from ACE file

```
mg_neutron_mode           //Process mode
ace_file_name      ( ../ace/H001.ace )       //ACE file name
ace_file_name_tsl  ( ../ace/01_h_in_h2o.ace ) //ACE file name (TSL data)
mg_edit_option      ( MATXS  GENDF )         //Output format
mg_file_name         H_in_H2O                 //Output file name
mg_structure ( xmas_nea-lanl_172 )           // ign=18 in GROUPR/NJOY
mg_weighting_spectrum ( 1/e )                //iwt=3 in GROUPR/NJOY
mg_tsl_data_type     ( hh2o )
```

## Input Example (5/7): Graphite

[illegible]

# Input Example (6/7): UO<sub>2</sub>

```

mg_neutron_mode                //Process mode
nucl_file_name ( ../lib/O016.dat ../lib/U235.dat ../lib/U238.dat )
mg_edit_option ( SimpleGENDF SimpleMATXS GENDF MATXS
  "1DXS 1, 2, 4, -50" "2DXS 1, 2, 4, -50" MGFlux )
  //"1DXS 1, 2, 4, -50": 1D cross-section data
  //"2DXS 1, 2, 4, -50": 2D cross-section data
  //MGFlux: Multi group flux data

mg_file_name mix_UO2           //Output file name
mg_structure ( xmas_nea-lanl_172 ) //ign=18 in GROUPR/NJOY
mg_weighting_spectrum ( 1/e )   // iwt=3 in GROUPR/NJOY
temp                600.0 //[K]
legendre_order      3

mg_number_density    //O16    U235    U238
                    (4.58e-2  7.18e-4  2.22e-2) //[1/barn/cm]

```

Atomic number density of each nuclide must be required to process compound of different isotopes.



# Input example (7/7): U-235

## (Automated setting of background XS)

```
mg_neutron_mode           //Process mode
nucl_file_name ( ../lib/U235.dat )    //ENDF file name
mg_file_name  U235              //Output file name
mg_edit_option ( GENDF  MATXS )
temp                300.0 //[K]
mg_structure ( ign18 )           //ign=18 in GROUPR/NJOY
mg_weighting_spectrum ( iwt03 )   // iwt=3 in GROUPR/NJOY
```

```
sigma_zero_data(auto 0.1 50 1.0e-10 rr linear)
```

```
//Tolerance = 0.1 (10%)
```

```
//Maximum number of background XS = 50
```

```
//Minimum background XS = 1.0E-10 (barn)
```

```
//Target of interpolation (factor / rr) = reaction rate)
```

```
//Interpolation method (cubic / linear) = linear interpolation
```

# IFENG (weight\_option) for TSL data generation

- Please check the IFENG value in the ACE file.
- IFENG=2 (weight\_option: tabulated) is strongly recommended to generate multi-group cross section file.
  - The prediction accuracy of transport calculation becomes lower when IFENG=0 or 1.

h-h <sub>2</sub> O,00t	0.999167	2.5852e-08	20220208
1001	0	0	0
0	0	0	0
0	0	0	0
0	0	0	0
11774842	3	31	80
0	0	0	0
1	500	998	0
0	0	0	0
0	0	0	0
0	0	0	0
498	1.0000000000e-11	1.0312500000e-11	1.0625000000e-11
1.0937500000e-11	1.1250000000e-11	1.1562500000e-11	1.1875000000e-11
1.2187500000e-11	1.2500000000e-11	1.2812500000e-11	1.3125000000e-11
1.3437500000e-11	1.3750000000e-11	1.4375000000e-11	1.5000000000e-11

IFENG

# NJOY compatible input format

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# Basics of NJOY compatible input format (1/2)

- Each module is independent.
  - RECONR, BROADR, GASPR, PURR, THERMR, ...
  - Input for each module is required.
- Order of module has impact on processing result.
  - RECONR must be set before BROADR.
  - BROADR must be set before PURR and THERMR.
- Comment line is after slash “/”.
  - Input information is written in NJOY manual and source files.
  - Input information uses Fortran language.
    - $A.gt.B \rightarrow A > B$ ,  $A.ge.B \rightarrow A \geq B$ ,  $A.lt.B \rightarrow A < B$ ,  $A.le.B \rightarrow A \leq B$
- Text data is surrounded by single quotation mark.
  - For example, ‘This PENDF file is U-235 from JENDL-4.0’.

# Basics of NJOY compatible input format (2/2)

- Card number of input information is line number.
  - The input parameters written in Card N must be set in one line.
  - If user sets slash, other input data are default values.
    - Third, forth, and fifth data use default values if five data are required in Card N and only two data are set.
  - NJOY needs MAT number to identify nuclide or material.
  - NJOY considers that data of many nuclides are written in one file.
  - NJOY can process data of several nuclides with one input file.
- “tapeNN” is only available for file name.
  - tape20, tape21, tape22, ..., tape99

# MODER module and binary tape file

- Many NJOY sample input files use MODER module before processing.
  - MODER module changes file format (ASCII, binary).
  - NJOY considers that this file is binary data if users set minus value.
    - Binary data is effective to reduce data size and readout time of files.
- FRENDY cannot handle binary data.
  - FRENDY considers that this file is ASCII data even if user sets minus value.

moder

*/ command*

20 -21

*/ nin, nout*

Original ENDF file name : tape20  
Changed file name (binary) : tape21

# NJOY modules handled in FRENDY

- FRENDY version 2 handles input files of the following NJOY modules.
  - MODER, RECONR, BROADR, GASPR, PURR, UNRESR, THERMR, ACER, GROUPE, and MATXS.
- Though THERMR input data of NJOY2012 and NJOY2016 are different from those of NJOY99, FRENDY can treat both input data.
  - FRENDY skips new input option “iform” in THERMR of NJOY2016.
  - FRENDY uses “iform=0: MF6 special”.
- Fast, thermal, and dosimetry (iopt=1~3) are only available for ACER module.
- FRENDY considers that this input file is PURR module even if user sets input file of UNRESR module.

# Notes on UNRESR and PURR modules

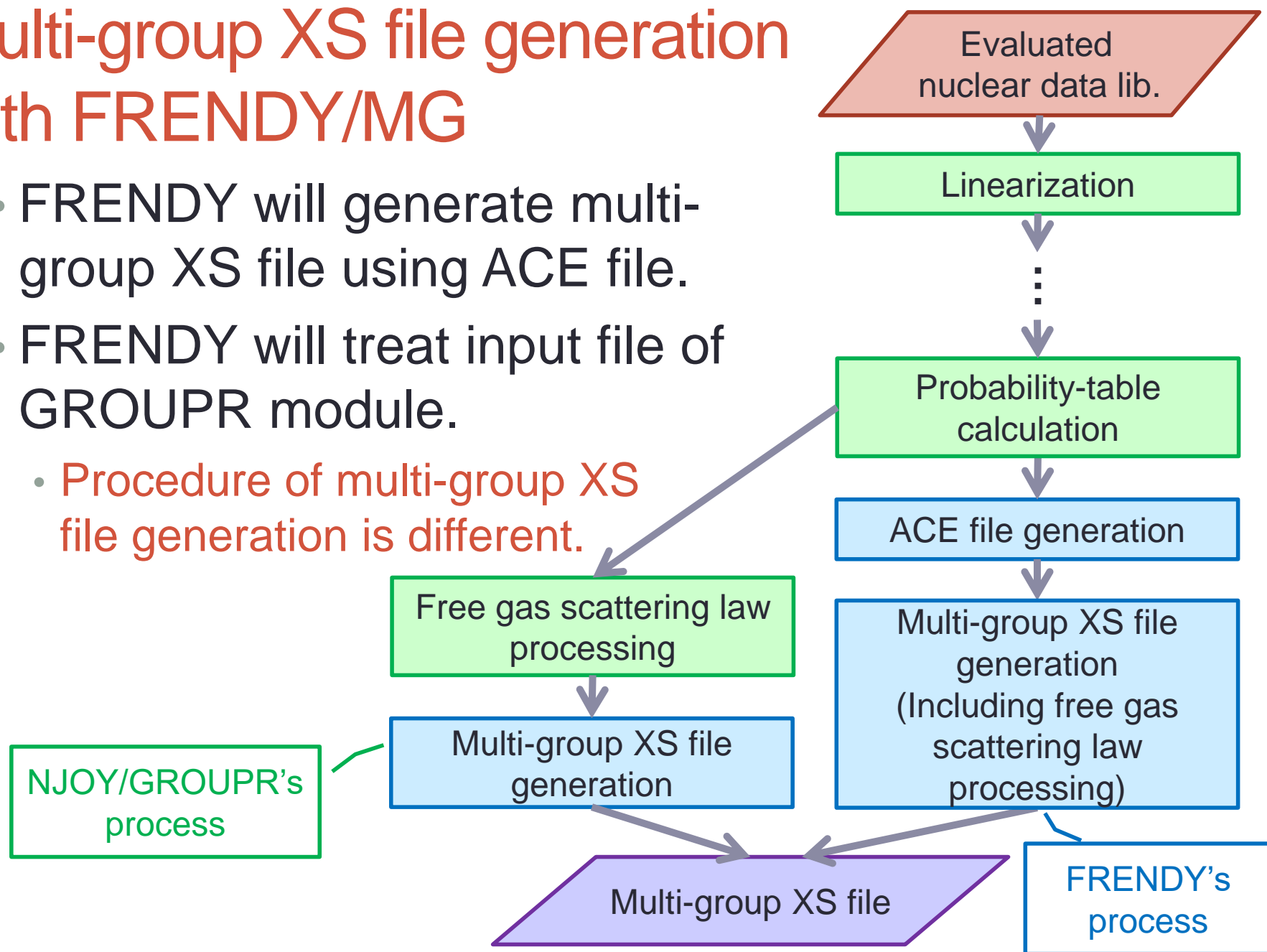
- Both modules treat self-shielding effect in the unresolved resonance region.
  - PURR is only available for continuous energy Monte Carlo codes.
  - PURR and UNRESR are available for multi-group calculation codes.
  - Processing time of UNRESR is much shorter than that of PURR.
- NJOY developers do not recommend to use UNRESR.
  - “For most purposes, UNRESR has been superseded by PURR.”\*)
  - “The PURR results may be more reliable at low  $\sigma_0$  values than UNRESR results.”\*)
- FRENDY doesn't have equivalent function of UNRESR.
  - FRENDY uses equivalent function to PURR module even if user uses input of UNRESR module.

\*) R. E. MacFarlane and A. C. Kahler, “Methods for Processing ENDF/B-VII with NJOY,” *Nucl. Data Sheets*, **111**, pp.2739-2890 (2010).



# Multi-group XS file generation with FRENDY/MG

- FRENDY will generate multi-group XS file using ACE file.
- FRENDY will treat input file of GROUPR module.
  - Procedure of multi-group XS file generation is different.



# NJOY compatible input file generation using FRENDY

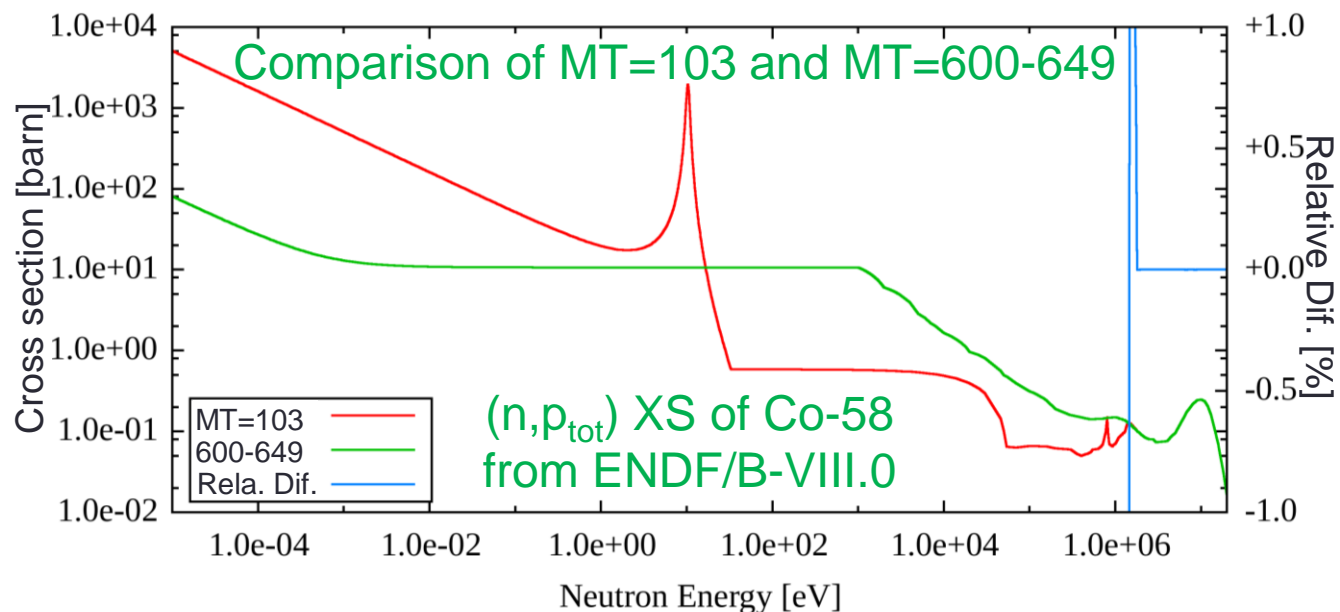
- FRENDY generates NJOY compatible input file using FRENDY input file.
  - Adding “\_make\_inp” at the end of processing mode
    - ace\_fast\_mode\_make\_inp, ace\_tsl\_mode\_make\_inp, ace\_dosi\_mode\_make\_inp
    - NJOY input file name is “FRENDY input file name”.njoy\_input.dat.
- We strongly recommend to check generated input file.
  - Major nuclides and materials in JENDL-4.0, ENDF/B-VIII.0, JEFF-3.3 have already checked.
  - Other materials, e.g.,  $\text{UO}_2$ ,  $\text{SiO}_2$ , ..., may be incorrect.
    - MAT number of TSL data depends on the evaluated nuclear data library.
    - FRENDY cannot distinguish material data from MAT number.

# Input check function of FRENDY

- Several input parameters largely affect processing results.
- FRENDY verifies processing results in each processing step.
  - E.g., linearization and processing of thermal scattering law data
  - This function is available for FRENDY input and NJOY compatible input.
  - This function will be helpful to make the correct input file.

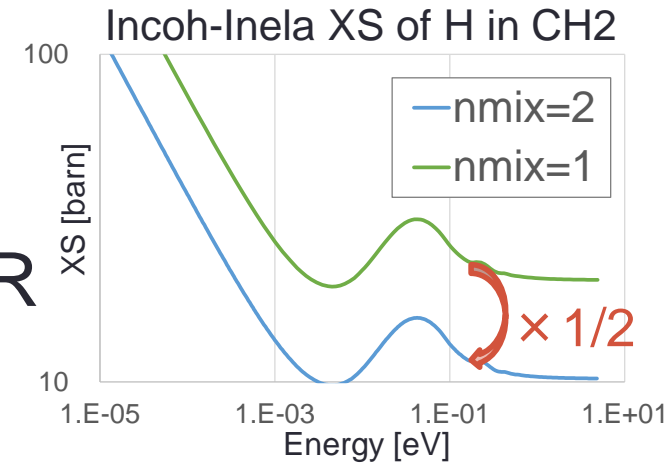
# Example of input check function (1/3)

- ENDF-6 format allows two types of particle production reaction (p, d, t, He-3,  $\alpha$ ).
- Total (MT=103-107) / each excited state (MT=600-849)
  - Total particle production cross section of some nuclides is not identical to sum of each excited state production cross section.
- FRENDY compares total and sum of each excited state XS
  - Output warning message if total XS is different to sum of each excited XS.



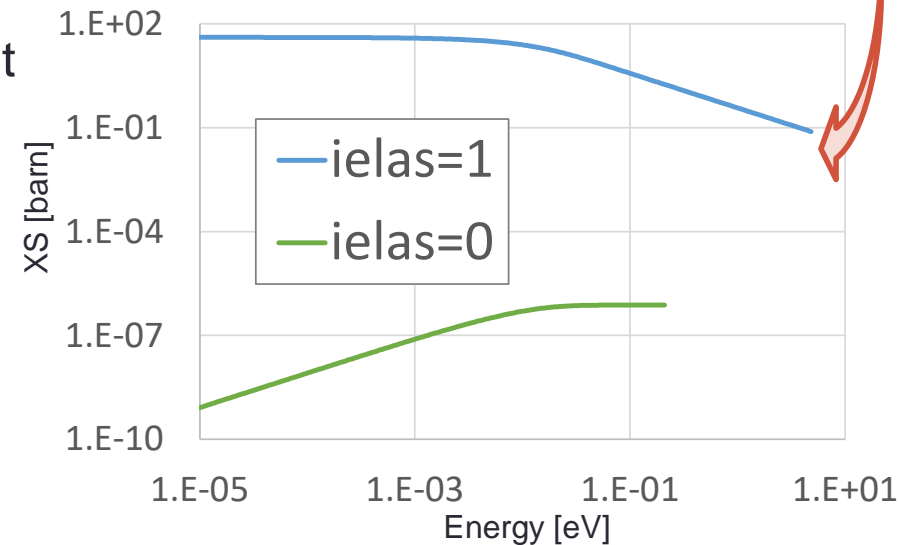
# Example of input check function (2/3)

- natom of THEMUR and nmix of ACER affect incoh-inela scattering XS.
  - natom: number of principal atoms
  - nmix: number of atom types in mixed moderator
    - These parameter modifies incoh-inela scattering XS.
- Users have to check XS to verify these parameters.
- **FRENDY automatically checks XS.**
  - Incoh-inela scattering XS is nearly equal to elastic scattering XS at the upper limit of incident energy.
  - FRENDY compares these XSs at the upper energy.
    - Output warning message if the relative difference is larger than 10%.



# Example of input check function (3/3)

- ACER requires elastic scattering type.
  - ielas of ACER : 0=coherent, 1=incoherent
- This parameter has large impact on processing results.
- FRENDY checks elastic scattering type using PENDF file.
  - LCT value of MF=6
    - LCT=7: coherent, LCT=6: incoherent
  - Output warning message if ielas of ACER is different to PENDF data.
- These input check functions will be helpful to make the correct input file.



# Concluding and remarks

- FRENDY treats two types of the input format.
- FRENDY input format
  - First line must be set processing mode.
    - ace\_fast\_mode, ace\_tsl\_mode, ace\_dosi\_mode
  - Other lines are free format.
    - Setting “input data name” and “input data”
    - Bracket is used to describe array data.
    - Text data is surrounded by single or double quotation mark.
    - Comment line is similar to C/C++.
- NJOY compatible input format
  - FRENDY version 1 handles input files of the following NJOY modules.
    - RECONR, BROADR, GASPR, PURR, UNRESR, THERMR, ACER, MODER.
  - Each module is independent
    - Input for each module is required
    - Order of module has impact on processing result.
  - Comment line is after slash
  - The input parameters written in Card N must be set in one line.
  - If user sets slash, other input data are default values
- Input checker function of FRENDY
  - FRENDY verifies processing results in each processing step.

# [Appendix]

## Explanation of NJOY input format

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# Input description: RECONR

- Card 1: nendf npend
  - ENDF (input) file name and PENDF (output) file name (20~99)
- Card 2: tlabel
  - Label (one line comment) of PENDF
- Card 3: mat ncards ngrid
  - MAT number, Number of comment lines of PENDF file, Number of additional energy grids
- Card 4: err tempr errmax errint
  - Allowable error of linearization, temperature [K], Maximum error of linearization, Allowable error of linearization (integral value)
- Card 5: cards
  - Comment line of PEND file (Line number = ncards)
  - ncards=0: Skip this card
- Card 6: enode
  - Additional energy grid (Grid number = ngrid)
  - grid=0: Skip this card

**Gray parameters:**  
**Recommended to**  
**use the default value.**

# Important point: RECONR

- nendf/npendf (card 1)
  - nendf and npendf are only available from 20 to 99.
  - File name of ENDF or PENDF is tapeXX (tape20~tape99).
    - ENDF file name is tape20 if nendf=20.
  - Original ENDF file name: nendf, Output PENDF file name: npendf
- err/errmax/errint (card 4)
  - We recommend to use  $\text{err} = 1.0\text{E-}3$  (0.1%).
  - Default values should be used for errmax and errint.
- tempr (card 4)
  - tempr must be set 0 K.
  - Single-Level Breit-Wigner resonance formula can directly calculate Doppler broadened XS using psi-chi approximation method.
    - The difference at top and bottom of resonance XS will become larger when psi-chi method is used.
  - Doppler broadened XS must be calculated in BROADR module.

# Example: RECONR

```
reconr           / command
20 21           / input(tape20), output(tape21)
'pendf tape for JENDL-4 U235' / identifier for PENDF
9228            / mat
1.00e-03        / err
0              /
```

ncards and ngrid  
are default value.  
(ncards/ngrid=0)

Last data is mat=0.  
(Card 3 of 2<sup>nd</sup> nuclide)

tempr, errmax, and  
errint are default value.

- NJOY considers that data of many nuclides are written in one file.
  - NJOY processes data of several nuclides with one input file if user repeats Card 3~Card 6.
  - **mat=0 in Card 3 means the end of RECONR module.**

# Input description: BROADR

- Card 1: nendf nin nout
  - ENDF file name, PENDF file name before BROADR (input PENDF), PENDF file name after BROADR (output PENDF)
- Card 2: mat1 ntemp2 istart istrap temp1
  - MAT number, number of temperatures, whether temp1 data is copy to nout or not, whether all temperature data is calculate from tmep1 or not, temperature of nin (normally 0 K)
- Card 3: errthn thnmax errmax errint
  - Allowable error of linearization, Maximum energy of Doppler broadening, Maximum error of linearization, Allowable error of linearization (integral value)
    - errthn, errmax, and erring of BROADR is identical to err, errmax, and errint of RECONR
- Card 4: temp2
  - Doppler broadened temperature (temperature number=ntemp2)

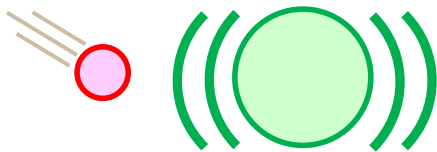
# Important point: BROADR

- thnmax (card 3)
  - NJOY considers that thnmax is the lowest of input value, upper boundary of resolved resonance regions, and 1 MeV.
  - Old version of NJOY, e.g., NJOY99 and NJOY2012, uses threshold energy as the thnmax.
    - U-235 in JENDL-4.0 cannot be correctly processed since threshold energies of several reactions are very small.
    - Minus thnmax value must be used to avoid this problem.
      - Input value is forcibly set when user sets minus thnmax value.
  - FRENDY and the current version of NJOY2016 do not check threshold energy.

# Threshold reaction and Doppler broadening

- Doppler broadening considers nucleus vibration.
- Threshold energy will also be moved by nucleus vibration.
  - Current nuclear data processing codes do not consider the moving of threshold energy.

[Reaction of incident particle and nucleus]

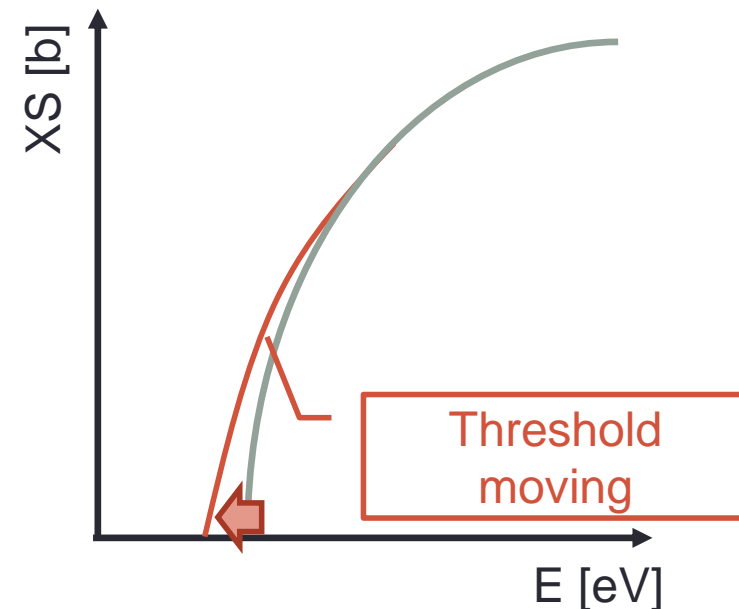


[Equation of Doppler broadening]

$$\sigma(v, T) = \frac{1}{v^2} \sqrt{\frac{\beta}{\pi}} \int_0^\infty dv_r v_r^2 \sigma(v_r) \left[ \frac{e^{-\beta(v-v_r)^2}}{-e^{-\beta(v+v_r)^2}} \right]$$

$T$ : Temperature,

$v$ : velocity of incident particle,  $v_r$ : relative velocity



# Example: BROADR

broadr

20 21 22

9228 1

1.00e-03 -5.0E+2

296.0

0

*/ command*

*/ endf, pendf(in), pendf(out)*

*/ mat, temp no*

*/ err, thnmax*

*/ temp*

*/*

Maximum energy is forced to 500 eV.

istart, istrap, and temp1 are default values.

Last data is mat=0.  
(Card 2 of 2<sup>nd</sup> nuclide)

errmax and errint are default values.

# Input description: GASPR

- Card 1 : nendf nin nout
  - ENDF file name, PENDF file name before GASPR (input PENDF), PENDF file name after GASPR (output PENDF)
- GASPR requires only ENDF/PENDF file names.
  - GASPR can be skipped if user does not need gas production XS (MT=203~207).

## [Example of GASPR]

gaspr	<i>/ command</i>
20 22 23	<i>/ endf, pendf(in), pendf(out)</i>



# Input description: PURR

- Card 1: `nendf nin nout`
  - ENDF file name, PENDF file name before PURR (input PENDF), PENDF file name after PURR (output PENDF)
- Card 2: `matd ntemp nsigz nbin nladr iprint nunx`
  - MAT number, number of temperatures, number of background XS, number of probability bins, number of ladders, output option, number of energy grids (0: all)
- Card 3: `temp`
  - temperature (temperature number = ntemp)
- Card 4: `sigz`
  - Background XS  $\sigma_0$  (sigz number = nsigz)

# Important point: PURR

- nbin/nladr (card 2)
  - nbin must be larger than or equal to 15.
  - Recommended nladr is larger than or equal to 100.
- iprint (card 2)
  - Probability table information is output in output file if iprint = 1.
- nunx (card 2)
  - Recommended nunx is 0, *i.e.*, processing all energy grids.
  - Considering to current computing performance, the processing time is not so long.
- sigz (card 4)
  - Back-ground XS data **must be descending order.**
  - The first back-ground XS data must set large value, *e.g.*, 1.0E+10.
    - NJOY considers that the first back-ground XS is infinite.
    - FRENDY sorts back-ground XS data.
    - FRENDY considers that the maximum back-ground XS is infinite.

# Example: PURR

purrr	/ command
20 23 24	/ endf, pendf(in), pendf(out)
9228 1 10 20 100 0	/ mat, temp no, sig no, bin no, lad no
296.0	/ temp
1E10 1E4 1E3 300 100 30 10 1 0.1 1.0E-5	/ sig zero
0	/

This value is considered as infinite.

nunx is default value.

Last data is mat=0.  
(Card 2 of 2<sup>nd</sup> nuclide)

# Input description: THERMR

- Card 1: nendf nin nout
  - ENDF file name of TSL data, PENDF file name before THERMR (input PENDF), PENDF file name after THERMR (output PENDF)
- Card 2: matde matdp nbin ntemp iin icoh iform natom  
mtref iprint
  - MAT number of TSL data, MAT number of nuclide, number of equi-probable angular bins, number of temperatures, inelastic scattering option, elastic scattering option, output format option, number of principal atoms, MT number of inelastic scattering (221~225 only), output option
- Card 3: temp
  - Temperature (temperature number = ntemp)
- Card 4: tol emax
  - Allowable error of linearization, maximum energy to consider thermal scattering
    - tol is identical to err in RECONR.

Green parameters:  
User have to read TSL  
data to set these  
parameters.

# Important point: THERMR (1/3)

- nendf/nin (card 1) & matde/matdp (card 2)
  - nendf and matde are obtained from TSL data.
    - For example, TSL data file name of HinH2O and MAT number
    - nendf and matde are 0 for free gas scattering law processing.
  - nin and matdp are obtained from nuclide data.
    - For example, H-001 for HinH2O.
- iform (card 2)
  - Data format of angular and energy distribution can be changed using this option.
  - This option is added in NJOY2012.
    - NJOY2012/2016 input is incompatible to NJOY99 input.
    - Users have to check whether this input file has iform or not.
    - FRENDY can treat both input format.
- mtref (card 2)
  - MT number of elastic scattering is mtref+1.

# Important point: THERMR (2/3)

- natom (card 2)
  - Number of target nuclides in a molecule.
    - For example, HinH<sub>2</sub>O: natom=2, OinH<sub>2</sub>O: natom=1
- temp (card 3)
  - Temperature is **only available for given temperatures in TSL data.**
  - PENDF file of target nuclide must be broadened in this temperature.
- emax (card 4)
  - Recommended value is the maximum energy of TSL or higher energy, e.g., 10 eV.
  - Short Collision Time (SCT) approximation is used if the energy is higher than the given energy range in TSL data.

# Important point: THERMR (3/3)

- iin (card 2)
  - Meaning of iin value of NJOY2012/2016 is different to NJOY99.
  - User has to check whether TSL data has inelastic scattering data in MF=7/MT=4 or not.
- icof (card 2)
  - User has to check whether TSL data has elastic scattering data in MF=7/MT=2 or not.
  - Elastic scattering XS of graphite is added in output PENDF file if user sets icof=2 and TSL data does not have elastic scattering data.
    - icof=2: compute elastic scattering XS using TSL data or elastic scattering XS of graphite
- iform (card 2)
  - This option is added in NJOY2012.
  - NJOY2012/2016 input is incompatible to NJOY99 input.
  - User have to check whether this input file has iform or not.

# iin/icof options of NJOY99 and NJOY2016

## NJOY99

### iinc inelastic options

- 0 none
- 1 compute as free gas
- 2 reserved
- 3 reserved
- 4 read s(a,b) and compute matrix

### icoh elastic options

- 0 none
- 1 compute using ENDF6 input tape data
- or for pre-ENDF6 input set icoh =
- 1 graphite
- 2 beryllium
- 3 beryllium oxide
- 11 polyethylene
- 12 h(zrh)
- 13 zr(zrh)

## NJOY2016

### iin inelastic options

- 0 none
- 1 compute as free gas
- 2 read s(a,b) and compute matrix

### icoh elastic options

- 0 none
- 1 compute using ENDF6 format data
- or for earlier formats
- 1 graphite
- 2 beryllium
- 3 beryllium oxide
- 11 polyethylene
- 12 h(zrh)
- 13 zr(zrh)

Option number is identical!!



# Example: THERMR

TSL data of HinH2O does not  
have elastic scattering data.  
(icoh = 0)

```
thermr
26 23 27
1 125 60 1 2 0 0 2 221
iform natom mtref
293.6
1.0e-03 1.00001e+01
```

*/command*  
*/nendf nin nout*  
*/matde matdp nbin ntemp iin icoh*  
*/tempr*  
*/tol emax*

Since there is iform option, this  
is NJOY2012/2016 input data.

# Input description: ACER (common)

- Card 1: nendf npend ngend nace ndir
  - ENDF file name, PENDF file name before ACER (input PENDF), multi-group photon data, ACE file name, XSDIR file name
- Card 2: iopt iprint itype suff nxtra
  - Processing mode, output option, ACE file mode (1=ASCII, 2=binary), suffix ID, number of IZ/AW data
    - **FRENDY can not handle itype=2.**
    - iopt=1: neutron/proton induced, iopt=2: TSL, iopt=3: dosimetry, iopt: 4: photo-atomic, iopt=5: photo-nuclear
- Card 3: hk
  - Label (one line comment) of ACE (Maximum: 70 characters)
- Card 4: iz, aw
  - IZ/AW pair data (number of iz, aw = nxtra)
  - nxtra=0: Skip this card

# Input description: ACER (iopt=1)

- Card 5: matd tempd
  - MAT number, temperature
- Card 6: newfor iopp
  - Whether new angular distribution format (LAW=61) is used or not, whether detailed photon data is used or not
    - Default newfor option is 1 (use LAW=61).
    - MCNP4C and later version of MCNP can handle LAW=61.
- Card 7: thin(1) thin(2) thin(3)
  - Data reduction option
    - User should not enter this option and should skip this card using slash “/”.

# Input description: ACER (iopt=2)

- Card 8: matd tempd tname
  - MAT number of nuclide, temperature,  $S(\alpha, \beta)$  identifier of MCNP, e.g., lwtr (Maximum: 6 characters)
- Card 8a: iza01 iza02 iza03
  - ZA value of target nuclide
- Card 9: mti nbint mte ielas nmix emax iwt
  - MT number of inelastic scattering, number of energy bins of 2<sup>nd</sup> particle, MT number of elastic, whether coherent elastic (ielas=0) or incoherent elastic (ielas=1), number of atom types in mixed moderator, maximum energy to consider thermal scattering, output format of energy of 2<sup>nd</sup> particle
    - mti is 0 or mtref of THERMR, mte is 0 or mtref+1 of THERMR
    - Emax is emax of THERMR
    - iwt=0: skewed, iwt=1:discrete, iwt=2:continuous
      - MCNP6.1 or later version can handle iwt=2.

# Important point: ACER (iopt=2)

- tname (card 8)
  - This name is also used in MCNP input file.
- iza01 iza02 iza03 (card 8a)
  - iza02 and iza03 are normally 0.
  - Target nuclide have several isotopes, e.g., Zr in ZrH, user has to set iza02 and iza03.
- ielas (card 9)
  - User has to check TSL data (MF=7/MT=2) and including data type.
- nmix (card 9)
  - Mainly, Benzene, BeO, and SiO<sub>2</sub> are nmix=2 and others are nmix=1
    - Though BeO from JENDL-4.0 is nmix=2, BeO from ENDF/B-VIII.0 is nmix=1.
    - **Please carefully check TSL data to process Benzene, BeO, and SiO<sub>2</sub>.**
- iwt (card 9)
  - MCNP6.1 or later version of MCNP can handle iwt=2.
  - Recommended iwt value is iwt=0 if user uses PHITS code or previous version of MCNP.
    - The nuclear calculation results using iwt=1 may no be incorrect.

# Example: ACER (iopt=1)

If user does not use photon data, ngend is 0.

acer	<i>/ command</i>
20 24 0 30 31	<i>/ nendf, npend, ngend, nace, ndir</i>
1 1 1 0.00	<i>/ iopt(fast), iprint(max), itype, suffix</i>
'ACE file for JENDL-4 U235'	<i>/ descriptive character</i>
9228 296.0	<i>/ mat, temp</i>
1	<i>/ newfor(yes)</i>
	<i>/ thin(1), thin(2), thin(3)</i>
stop	<i>/</i>

Skip Card 7 (thin(1)~thin(3)).

# Example: ACER (iopt=2)

acer	<i>/command</i>
26 27 0 30 31	<i>/nendf npend ngend nace ndir</i>
2 1 1 0.0 0	<i>/iopt iprint itype suff nxtra</i>
'ACE file for HinH2O from JENDL-4.0'	<i>/hk</i>
125 2.93600e+02 'lwtr'	<i>/matd tempd tname</i>
1001	<i>/iza</i>
221 60 0 0 1 1.00001e+01 0	<i>/mti nbint mte ielas nmix emax iwt</i>

User considers that 'lwtr' means HinH2O.  
(User has to use 'lwtr' in MCNP input file.)