

Operator DYNA_SPEC_MODAL

1 Drank

Compute the response by modal recombination of a linear structure with a random excitation. This excitation is defined in the form of interspectrums of power of modal forces. The response is drawn up in the form of interspectrums of power of generalized displacements.

Each interspectral matrix is associated with a modal base for which the response is calculated. Result the product is a concept of type `interspectrum`.

2 Syntax

```
tinsp [interspectrum] = DYNA_SPEC_MODAL (

◆BASE_ELAS_FLUI=bef , [melasflu]
◆VITE_FLUI=vitefl [R]
◇PRECISION=/ 1.E-3 [DEFAULT]
/prec [R]
◆EXCIT=_F (
◆INTE_SPEC_GENE=exc , [interspectrum]
)

◇OPTION=/ "TOUT" [DEFAULT]
/ "DIAG",

◇TITER=titer , [TXM]

);
```

3 Operands

3.1 Operand **BASE_ELAS_FLUI**

◆ `BASE_ELAS_FLUI = bef`

Concept of the `melasflu` type, defines a set of modal bases associated with the various rates of flow with the fluid.

3.2 Key word **VITE_FLUI**

◆ `VITE_FLUI = vitefl`

Rate of flow of the fluid for the computation of response.

3.3 Key word **accuracy**

◇ `accuracy = prec`

Accuracy on the rate of flow of the fluid (by default 1 E-3).

3.4 Key word **EXCIT**

◆ **EXCIT**

Key word factor which defines the excitation.

◆ `INTE_SPEC_GENE = exc`

Concept of type `interspectrum`, defines the interspectrums of excitation (modal forces).

3.5 Operand **OPTION**

◇ `OPTION = "TOUT" or "DIAG"`

Argument of type text which indicates if one wants to calculate all the interspectrums "TOUT" or autospectrums only "DIAG". By default one calculates all the interspectrums.

3.6 Operand **TITER**

◇ `TITER = title`

Argument of type text defining the title attached to the concept `interspectrum` in output.