

Operator MACR_ELEM_DYNA

1 Goal

To define a dynamic macronutrient of under-structuring.

Within the framework of a transitory, modal or harmonic analysis, with dynamic under-structuring the operator `MACR_ELEM_DYNA` carry out the projection of the matrices of rigidity, mass and possibly of damping (analyzes harmonic) on the basis of modal substructure defined by `DEFI_BASE_MODAL` [U4.64.02], and the extraction of the matrices of connection of the interfaces. The result is consisted by the projected matrices and of the matrices of connection. It can be used several times with different orientations in the same model (cf. `DEFI_MODELE_GENE` [U4.65.02]). It can be printed on file by the order `IMPR_MACR_ELEM` [U7.04.33].

Product a concept of the type `macr_elem_dyna`.

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2 Syntax

```
macro_dyna [macr_elem_dyna] = MACR_ELEM_DYNA

(
  ◆ BASE_MODAL = bamo, [mode_meca]
  ◇ MACR_ELEM_DYNA = macro_dyna, [macr_elem_dyna]
)

# Given matrices :

◇ / MATR_RIGI = Mr., [matr_asse_DEPL_R]
[matr_asse_DEPL_C]

/ MATR_MASS = mm, [matr_asse_DEPL_R]

/ MATR_IMPE = semi, [matr_asse_gene_C]

# If well informed MATR_IMPE:
◆ FREQ_EXTR = freq, [R]
◇ AMOR_SOL = / 0.0, [DEFECT]
/ amosol, [R]
◇ MATR_IMPE_INIT = mi0, [matr_asse_gene_C]
/ | MATR_IMPE_RIGI = Mr., [matr_asse_gene_C]
| MATR_IMPE_AMOR = my, [matr_asse_gene_C]
| MATR_IMPE_MASS = mm, [matr_asse_gene_C]

◇ / MATR_AMOR = my, [matr_asse_DEPL_R]
/ AMOR_REDUIT = , [l_R]

◇ / GROUP_NO = grno, [group_no]
/ SANS_GROUP_NO = grno, [group_no]

# Under-structuring static:

◇ CAS_CHARGE = _F (
  ◆ NOM_CAS = nocas , [k8]
  ◆ / VECT_ASSE_GENE= vgen , [vect_asse_gene]
  / RESU_GENE= resugen , [tran_gene]
),

# manual Filling of the reduced matrices (given experimental):

◇ MODELE_MESURE = _F (
  ◆ FREQ = freq , [l_R]
  ◆ MASS_GENE = mgen , [l_R]
  ◇ AMOR_REDUIT = xsi, [l_R]
),

)
```

3 Operands

3.1 Operand BASE_MODAL

◇ BASE_MODAL = bamo

Name of the concept `mode_meca` product by the operator `DEFI_BASE_MODAL` [U4.64.02].

3.2 Operand MYCR_ELEM_DYNA

◇ MYCR_ELEM_DYNA = macro_dyna

Name of the concept mydynamic cro-element of type `myCr_elem_dyna` identical in the name of the produced concept. It is thus D-entering when it is present. The contents of the concept are modified starting from the data of the operands `MATR_IMPE*` present.

3.3 Operand MATR_RIGI

◇ MATR_RIGI = Mr.

Name of the concept stamps assembled of type `matr_asse_DEPL_R` or `matr_asse_DEPL_C` product by the operator `ASSE_MATRICE` [U4.61.22] or the macro-order `ASSEMBLY` [U4.61.21] corresponding to the matrix of rigidity of the substructure.

3.4 Operand MATR_MASS

◇ MATR_MASS = mm

Name of the concept stamps assembled of type `matr_asse_DEPL_R` product by the operator `ASSE_MATRICE` [U4.61.22] or the macro-order `ASSEMBLY` [U4.61.21] corresponding to the matrix of mass.

These two operands are to be employed if the modal base is used `bamo` is of type 'RITZ'.

3.5 Operand MATR_AMOR / AMOR_REDUIT

◇ / MATR_AMOR = my

Name of the concept stamps assembled of type `matr_asse_DEPL_R` product by the operator `ASSE_MATRICE` [U4.61.22] or the macro-order `ASSEMBLY` [U4.61.21] corresponding to the matrix of damping viscous, specific to the macronutrient. This damping must be of RAYLEIGH type by element (linear combination of rigidity and the mass on the level of the element) and is thus defined by the properties of the material (operator: `DEFI_MATERIAU` [U4.43.01], operands `AMOR_ALPHA` and `AMOR_BETA`).

/ AMOR_REDUIT =

List of reduced depreciation (percentage of damping criticizes) correspondent with each mode of vibration of the macronutrient. The length of the list is (with more) equal to the number of clean modes of the modal base; if it is lower, one supplements the list with reduced depreciation equal to the last term of the list entered by the user. No damping is associated with the static modes. The matrix of damping generalized of the macronutrient \bar{c}^k is thus diagonal incomplete (j index of the clean mode):

$$\bar{c}^k = \begin{pmatrix} \xi_j & 0 \\ 0 & 0 \end{pmatrix}$$

3.6 Operands MATR_IMPE / FREQ_EXTR / AMOR_SOL

◇ MATR_IMPE = semi

Name of the concept stamps assembled of type `matr_asse_gene_C` product by the operator `LIRE_IMPE_MISS` [U7.02.32] corresponding to the matrix of impedance of ground constitutive of the macronutrient.

◆ `FREQ_EXTR = freq`

Frequency of extraction of the matrix of impedance of ground necessary for the calculation of the matrix of radiative damping of ground starting from the imaginary part of the matrix `mi`.

◇ `AMOR_SOL = amosol`

Value of damping reduces material ground. It serves to distinguish in damping as the ground the properly material part and the radiative part. If it is nonnull, the radiative part `C` express yourself then such as:

$$2\pi \text{freq} C = \text{Imag}(mi(\text{freq})) - 2 \text{amosol} \text{Reel}(mi(\text{freq}))$$

3.7 Operand `MATR_IMPE_INIT`

◇ `MATR_IMPE_INIT = mi0`

Name of the concept stamps assembled of type `matr_asse_gene_C` product by the operator `LIRE_IMPE_MISS` [U7.02.32] correspondent with a matrix of impedance of ground constitutive of the macronutrient extracted at a quasi-worthless frequency. In particular in the cases of interaction ground-structure-fluid with the keyword `ISSF=' OUI'` in the call to `LIRE_IMPE_MISS`, that makes it possible to extract a contribution from mass `M` such as:

$$(2\pi \text{freq})^2 M = \text{Reel}(mi0) - \text{Reel}(mi(\text{freq}))$$

3.8 Operands `MATR_IMPE_RIGI/MATR_IMPE_AMOR/MATR_IMPE_MASS`

| `MATR_IMPE_RIGI = Mr.`
| `MATR_IMPE_AMOR = my`
| `MATR_IMPE_MASS = mm`

Name of the concepts of assembled matrix of type `matr_asse_gene_C` products by successive calls to the operator `LIRE_IMPE_MISS` [U7.02.32] in order to extract the respective contributions constitutive of the macronutrient in rigidity, damping or mass of a matrix of temporal impedance of ground. If at least of the operands is indicated, without others being present, then the contributions of the latter under the macronutrient are filled and put at 0.

An example of use is provided by the test `MISS03B` [V1.10.122].

3.9 OperandS `GROUP_NO/SANS_GROUP_NO`

◇ `GROUP_NO = grno`

Name of the group of nodes including the list of the nodes attaches with the degrees of freedom of the modes of interface by a relation `LIAISON_INTERF` (or `LIAISON_SOLIDE` if the dynamic interface is reduced to a node) with the nodes physical interface of the part of model on which one calculates the dynamic macronutrient. Its data is necessary only if this macronutrient is used as super-mesh of substructures defined by the keyword `AFFE_SOUS_STRUC` in a mixed model also including classical finite elements, and in this case, only when nodes of the interfaces physics and dynamics (the latter defined by `DEFI_INTERF_DYNA`) do not coincide. For example in the case of the dynamic interface reduced to a node connected by a solid connection to the physical interface.

◇ `SANS_GROUP_NO = grno`

Name of the group of nodes including the list of the nodes of the physical interface of the part of model on which one calculates the dynamic macronutrient. These nodes are in direct relationship to the nodes attached to the degrees of freedom of the modes of interface by a relation `LIAISON_INTERF` (or `LIAISON_SOLIDE` if the dynamic interface is reduced to a node). Its data is

necessary only if this macronutrient is used as super-mesh of substructures defined by the keyword `AFFE_SOUS_STRUC` in a mixed model also including classical finite elements, and in this case, only when nodes of the interfaces physics and dynamics (the latter defined by `DEFI_INTERF_DYNA`) do not coincide. For example in the case of the dynamic interface reduced to a node connected by a solid connection to the physical interface.

3.10 Keyword `CAS_CHARGE`

◇ `CAS_CHARGE`

This keyword factor makes it possible to define a set of loading cases **named** (keyword `NOM_CAS`). These loading cases are used to apply generalized vectors of load applied to the part of model on which one calculates the dynamic macronutrient so then this macronutrient is used as super-mesh of substructures in a mixed model also including classical finite elements.

3.10.1 Operand `NOM_CAS`

◆ `NOM_CAS = nocas`

The loading condensed under the name `nocas` (between “quotes”) corresponds to the loading defined by the argument `VECT_ASSE_GENE` or `RESU_GENE` on the part of model on which one calculates the dynamic macronutrient.

3.10.2 OperandS `VECT_ASSE_GENE/ RESU_GENE`

◆ `VECT_ASSE_GENE = vgen`

The loading condensed under the name `nocas` (between “quotes”) corresponds to the loading defined by Lbe argumentS alternate `VECT_ASSE_GENE` or `RESU_GENE`. It is obtained by projection that is to say of one vector assembled of load, maybe of a transitory result of force second member, applied to the part of model on which one calculates the dynamic macronutrient, on the modal basis `bamo` defined higher. These two options are tested simultaneously in the test `SDNX101B`.

3.11 Operand `MODELE_MESURE`

◇ `MODELE_MESURE`

This keyword factor makes it possible to manually fill the reduced matrices of the macronutrient, while using, for example, of the data resulting from measurements (and imported with `LIRE_RESU`). One must, has minimum, to return the generalized mass and the Eigen frequencies. One can also inform the list of reduced depreciation.

The well informed number of data must be equal to the number of modes of the modal base on which the macronutrient is built.

Not methodological: this kind of use of `MACR_ELEM_DYNA` justifies itself for the use of the method of structural modification starting from an experimental model. A presentation of the method is given in U2.07.03. **The modal base used to build the macronutrient should be made up only of the clean modes of the measured structure**, and does not have to comprise the static statements with the interface, because those are false (because not measured and, in the actual position of knowledge, nonmeasurable).

The cas-test `sdll137e` is an example of the implementation of methodology.

3.11.1 Operand `FREQ`

◆ `FREQ = freq`

List of the identified Eigen frequencies.

3.11.2 Operand **MASS_GENE**

- ◆ `MASS_GENE = farmhouse`

List of the identified generalized masses.

3.11.3 Operand **AMOR_REDUIT**

- ◆ `AMOR_REDUIT = xsi`

List of reduced depreciation identified.

4 Example

An example of use of this operator is given in the documentation of the operator `DEFI_SQUELETTE` [U4.24.01].