

# FORWARD-MODE ENZYME IN DEVELOPING CONSTITUTIVE MODELS

WITH RATEL

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University of Colorado Boulder

SIAM CSE23  
Feb 28, 2023

# DIFFERENTIATION IN SOLID MECHANICS

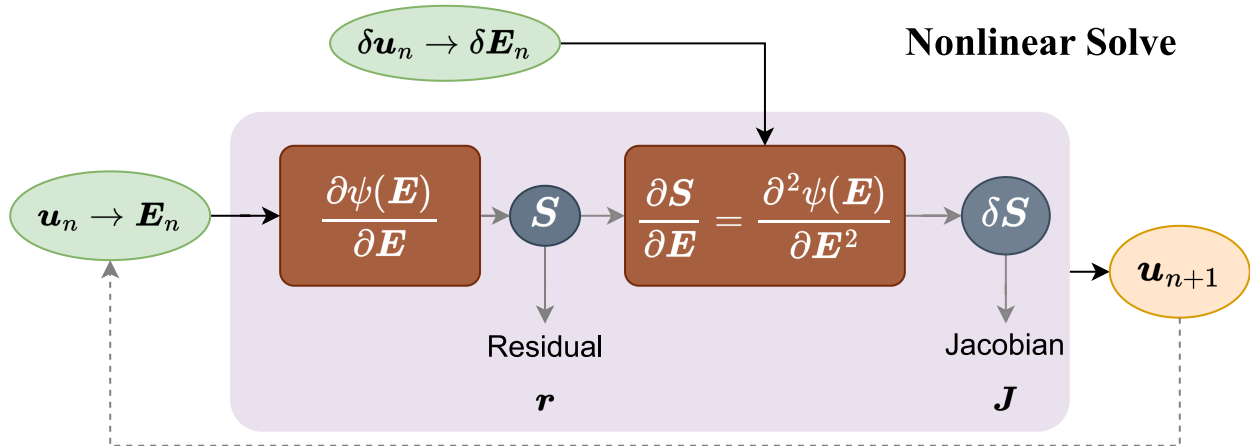
## FREE ENERGY FUNCTIONAL AND PDE SOLVERS

$$\psi(\mathbf{E}) = \frac{\lambda}{4} (J^2 - 1 - 2 \log J) - \mu (\log J + \text{trace } \mathbf{E}), \quad J = \sqrt{|\mathbf{I} + 2\mathbf{E}|}$$

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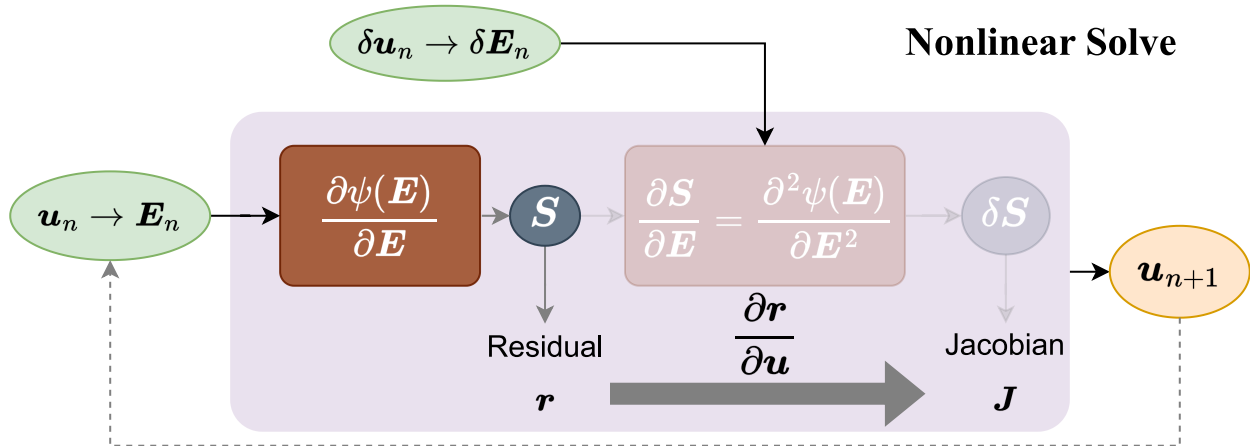
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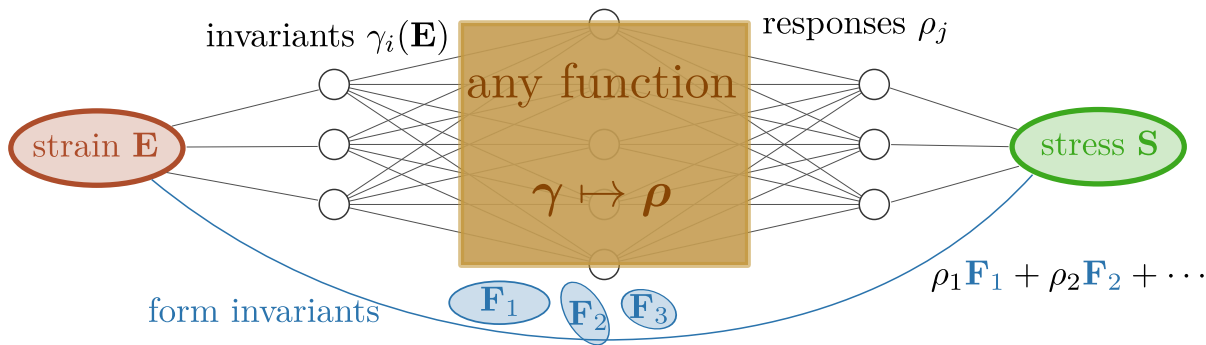
## FREE ENERGY FUNCTIONAL AND INVERSE PROBLEMS

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# ABAQUS

## UHYPER: USER SUBROUTINE

### Variables to be defined

U (1)

U, strain energy density function. For a compressible material, at least one derivative involving J should be nonzero. For an incompressible material, all derivatives involving J will be ignored. The strain invariants— $\bar{I}_1$ ,  $\bar{I}_2$ , and J—are defined in [Hyperelastic behavior of rubberlike materials](#).

U (2)

$\tilde{U}_{dev}$ , the deviatoric part of the strain energy density of the primary material response. This quantity is needed only if the current material definition also includes Mullins effect (see [Mullins effect](#)).

UI1 (1)

$$\partial U / \partial \bar{I}_1.$$

UI1 (2)

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UI1 (3)

$$\partial U / \partial J.$$

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- Fully automated commercial package (Solid Mechanics, FEM)

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- ▶ Fully automated commercial package (Solid Mechanics, FEM)
- ▶ Complex interface (too many inputs)
- ▶ Unstable for small deformation due to the choice of interface design

$$F = I + \nabla_X u$$

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$$F = I + \nabla_X u$$
- ▶ Not easy to change the interface

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AD helps us to create a more generic interface (one input function,  $\psi(E)$ ).

- Fully automated commercial package (Solid Mechanics, FEM)
- Complex interface (many inputs)
  - Not suitable for small amount of information due to the complexity of interface design
  - $\mathbf{F} = \mathbf{I} + \nabla_X \mathbf{u}$
- Not easy to change the interface

# RATEL: EXTENSIBLE, PERFORMANCE-PORTABLE SOLID MECHANICS

[HTTPS://GITLAB.COM/MICROMORPH/RATEL](https://gitlab.com/micromorph/ratel)

GitLab-CI passed

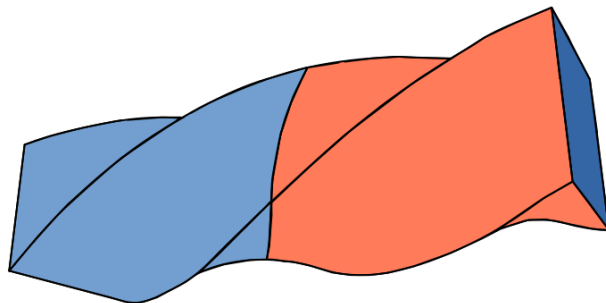
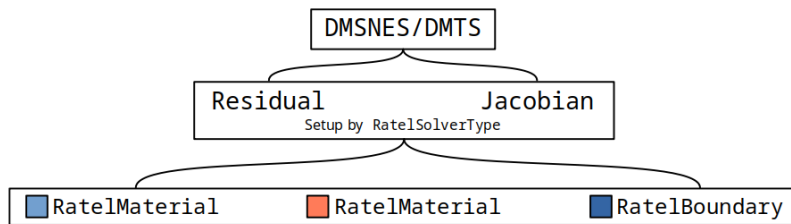
License BSD 2-Clause

Documentation latest

coverage 96.05%

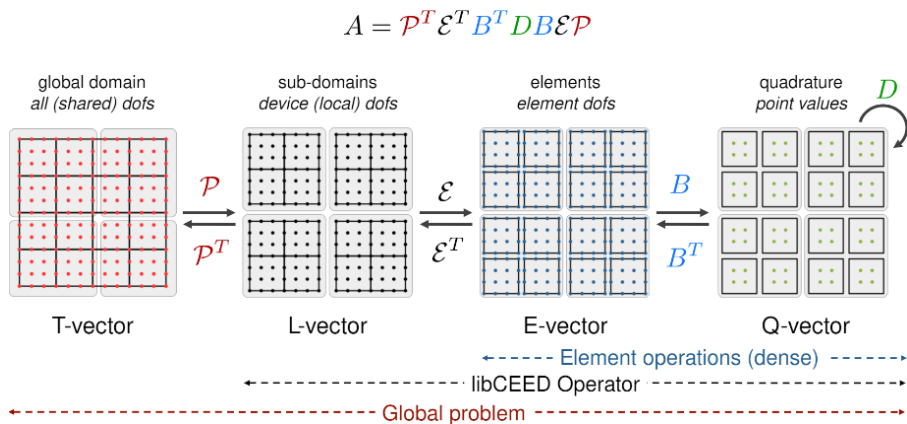
## Features:

- ▶ Linear elasticity
- ▶ Neo-Hookean and Mooney-Rivlin Hyperelasticity
- ▶ Multi-material
- ▶ Static, Quasistatic, Dynamic
- ▶ Initial and Current configurations



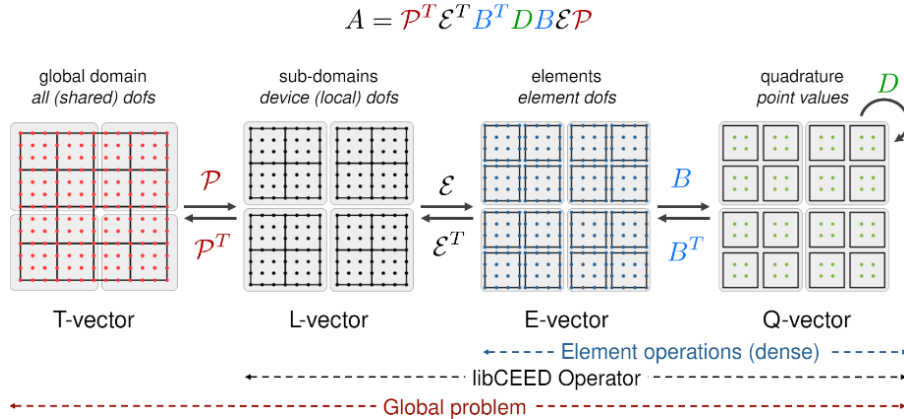
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COMPOSITION AND ABSTRACTION - LIBCEED: [HTTPS://LIBCEED.ORG/EN/LATEST/](https://libceed.org/en/latest/)



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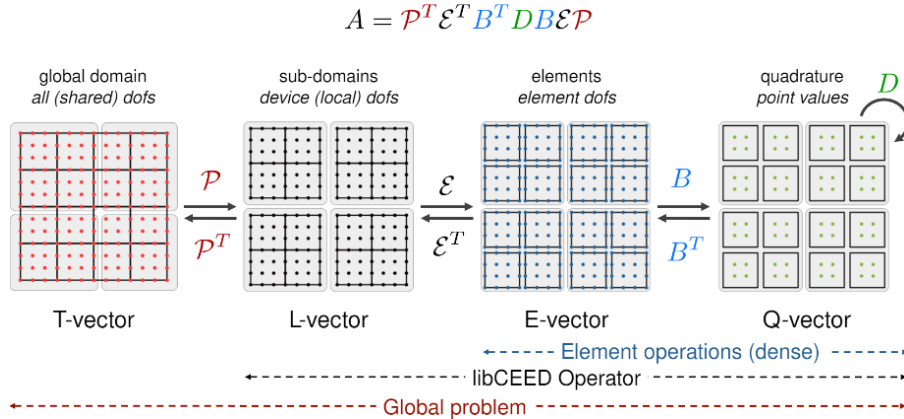
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## ► Purely algebraic high-order FEM

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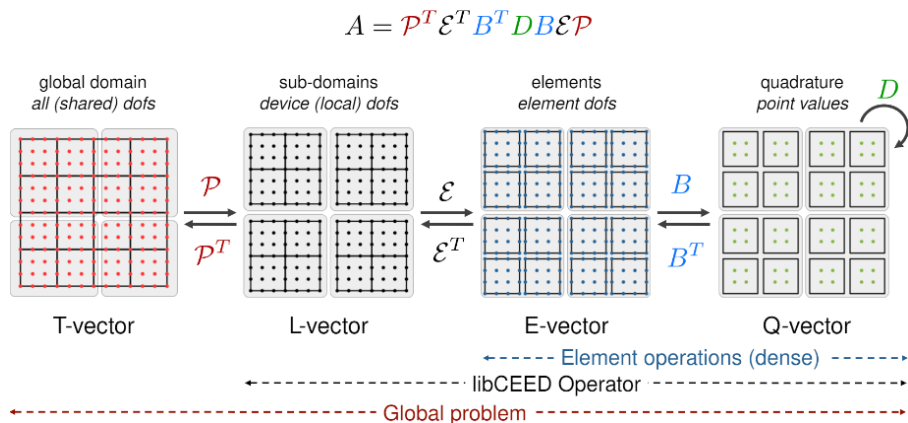
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- Purely algebraic high-order FEM
- Single source Vanilla C for physics

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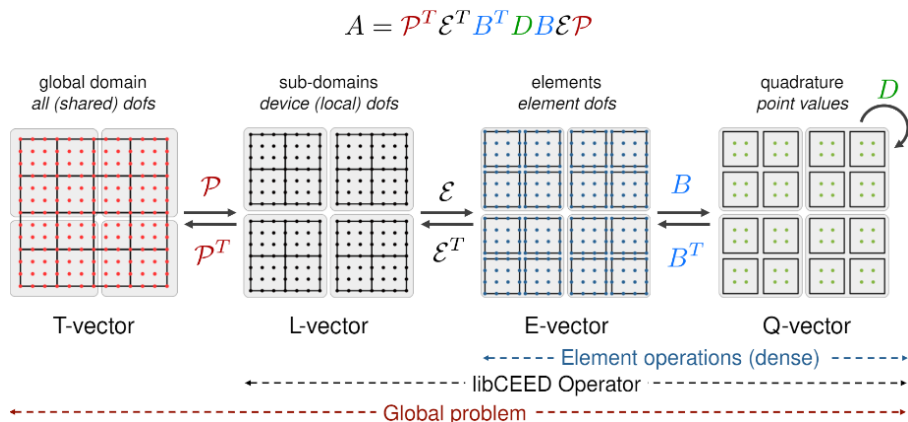


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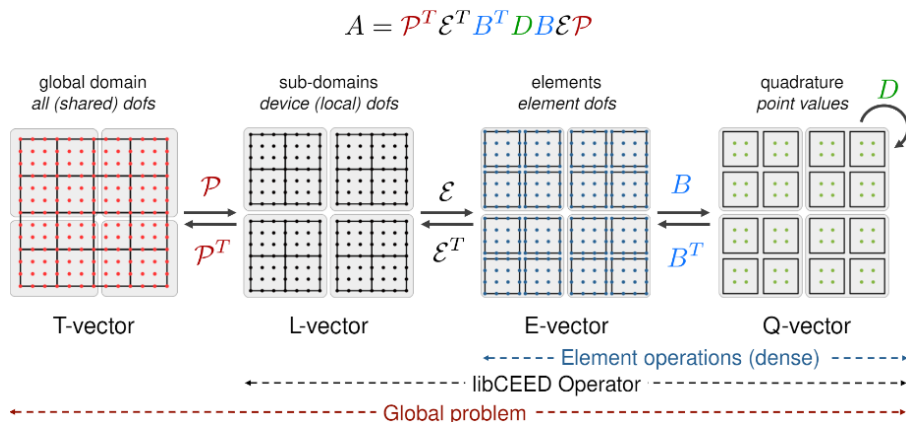
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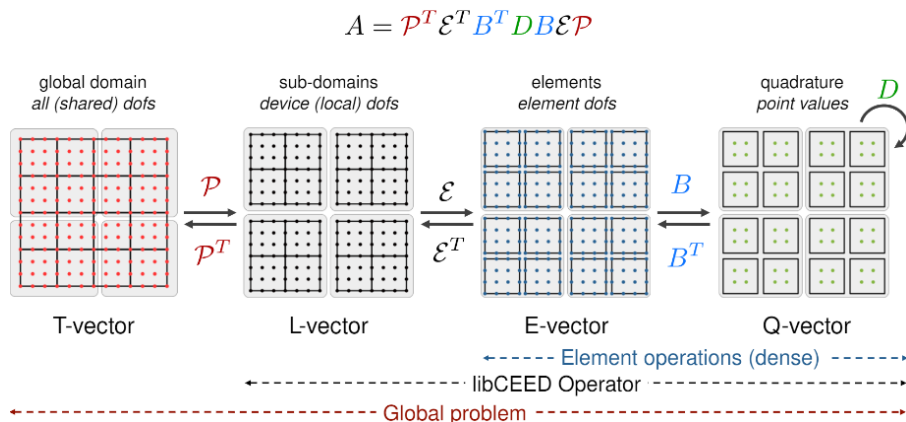
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- ▶ Support for Matrix-assembly and Matrix-free

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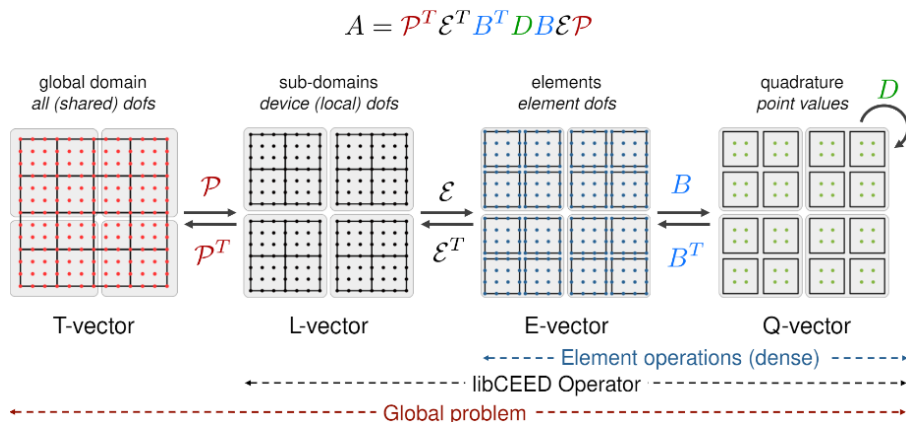
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- ▶ Support for Matrix-assembly and Matrix-free
- ▶ Operator abstraction
- ▶ User choice of data storage at quadrature point

# RATEL: EXTENSIBLE, PERFORMANCE-PORTABLE SOLID MECHANICS

## COMPOSITION AND ABSTRACTION - PETSc

### PETSc:

<https://petsc.org/release/>

- ▶ Parallel solution of PDEs

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- ▶ GPUs (CUDA, HIP, and OpenCL)

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- ▶ Optimization (PETSc/Tao)

# RATEL: EXTENSIBLE, PERFORMANCE-PORTABLE SOLID MECHANICS COMPOSITION AND ABSTRACTION - ENZYME AD

## Enzyme AD:

<https://enzyme.mit.edu/>

- ▶ High-Performance Automatic Differentiation

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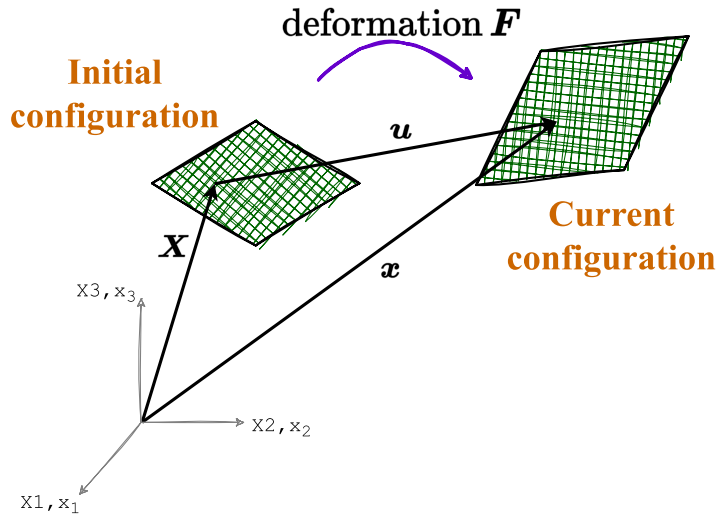
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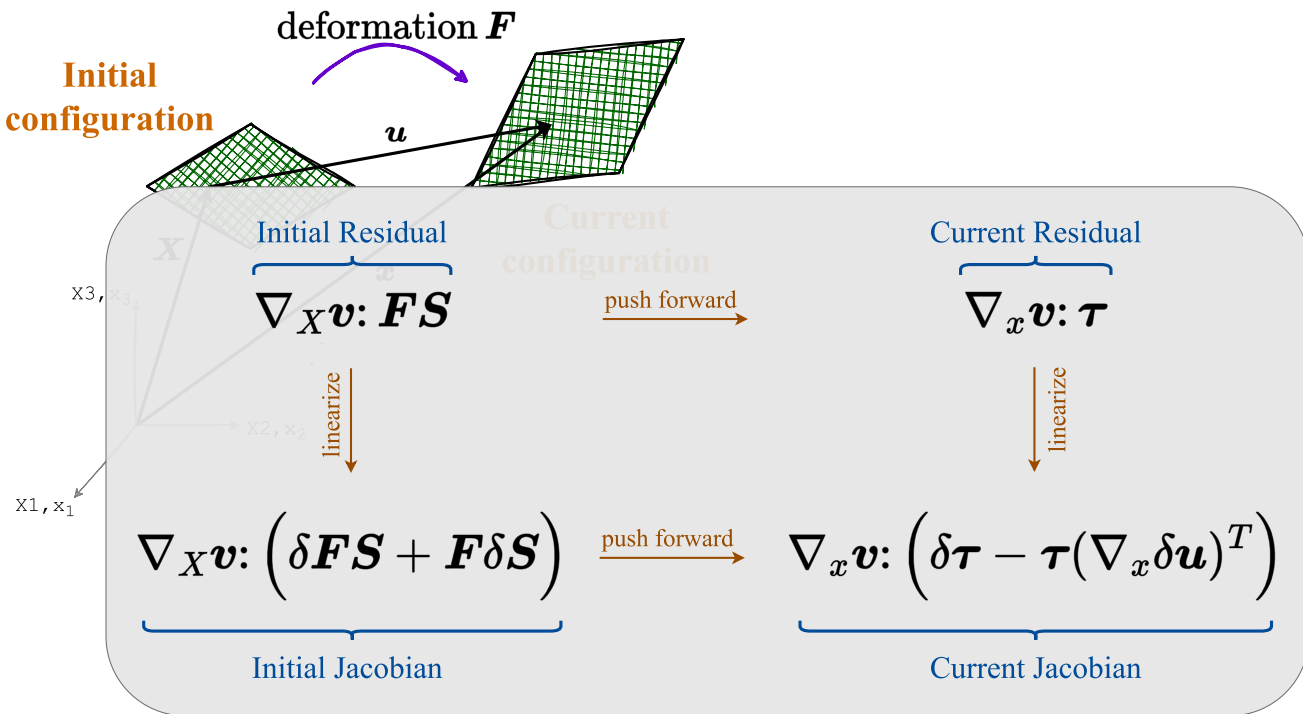
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- ▶ High-Performance Automatic Differentiation
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- ▶ Support for variety of languages (C/C++, Julia, Rust, Fortran, etc)
- ▶ *reverse* and *forward* mode AD

## INITIAL VS CURRENT CONFIGURATION



# INITIAL VS CURRENT CONFIGURATION



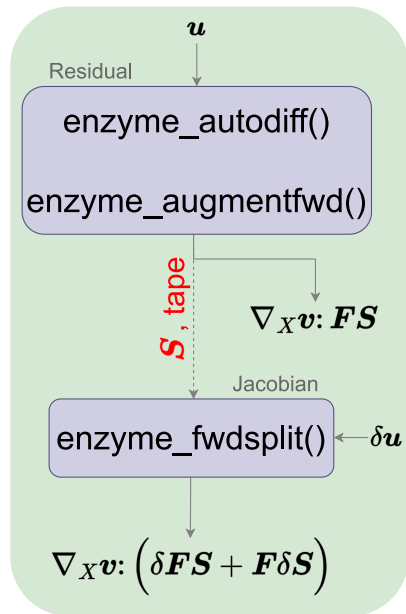
# RATEL - ENZYME AD

## INITIAL CONFIGURATION - REVERSE AND FORWARD SPLIT

```
// S = d(\psi) / d(E) [Reverse mode]
void SecondPiolaKirchhoffStress_NeoHookean_AD(...) {
    __enzyme_autodiff((void *)StrainEnergy, ...);
}

// Call forward S and return tape
__enzyme_augmentfwd(
    (void *)SecondPiolaKirchhoffStress_NeoHookean_AD,
    enzyme_allocated, tape_bytes, enzyme_tape, tape,
    enzyme_nofree, ...);

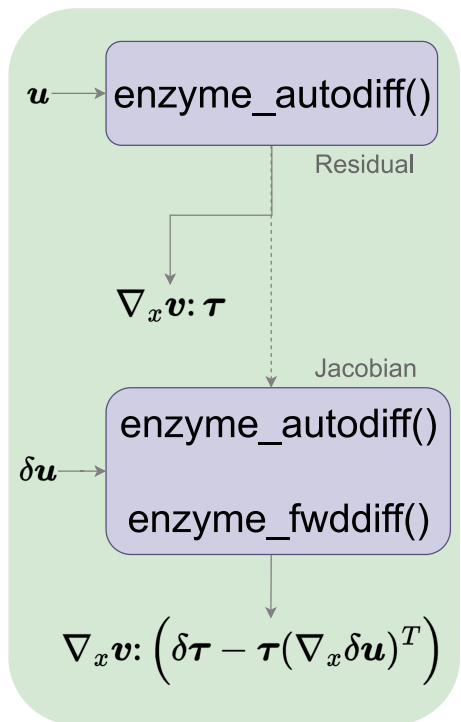
// Compute dS using the stored tape [Forward-split]
__enzyme_fwdsplit(
    (void *)SecondPiolaKirchhoffStress_NeoHookean_AD,
    enzyme_allocated, tape_bytes, enzyme_tape, tape, ...);
```





# RATEL - ENZYME AD

## CURRENT CONFIGURATION - REVERSE AND FORWARD



```
// Compute tau = (dPsi / de) * (2 e + I) [Reverse]
void Kirchhofftau_Voigt_NeoHookean_AD(...) {
    __enzyme_autodiff((void *)StrainEnergy, ...);
    ...
    for (int j = 0; j < 6; j++)
        b_Voigt[j] = 2 * e_Voigt[j] + (j < 3);
    ...
    RatelMatMatMult(1., dPsi, b, tau);
}

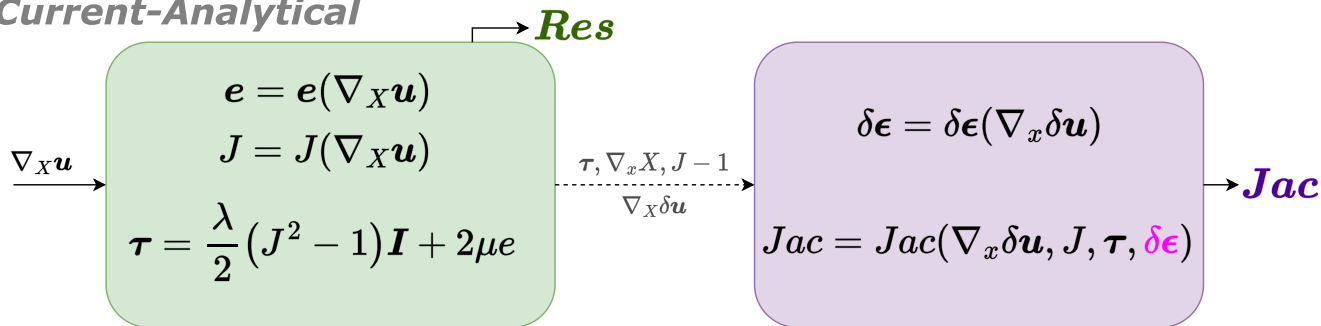
// Compute dtau [Forward]
CEED_QFUNCTION_HELPER void dtau_fwd(...) {
    __enzyme_fwddiff(
        (void *)Kirchhofftau_Voigt_NeoHookean_AD, ...);
}
```

## RATEL - ENZYME AD

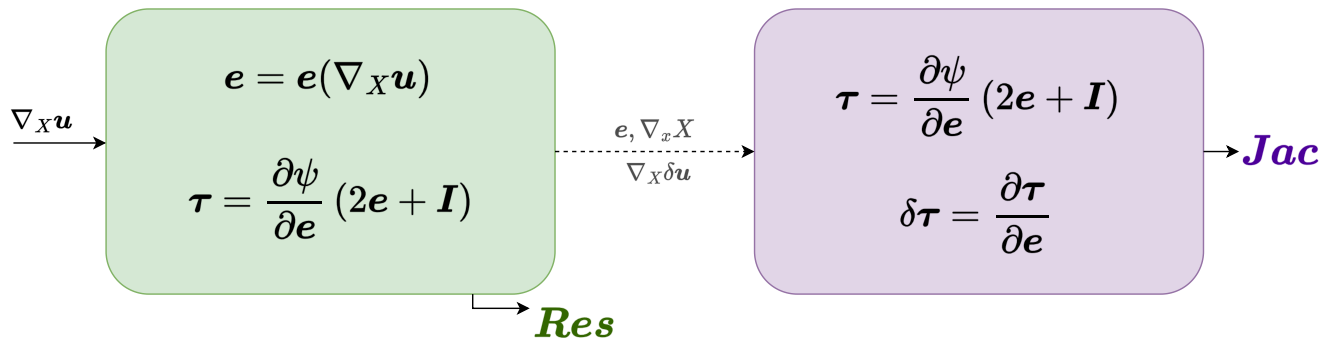
### PERFORMANCE FOR DIFFERENT JACOBIAN REPRESENTATIONS

Problem	Storage	Scalars	Time (s)
current	$W; \nabla_x \xi, \tau, J - 1$	17	36.2
initial	$W, \nabla_x \xi; \nabla_x u$	19	48.4
initial-AD	$W, \nabla_x \xi; \nabla_x u, S, \text{tape}$	31	53.9
current-AD	$W; \nabla_x \xi, e$	16	55.8

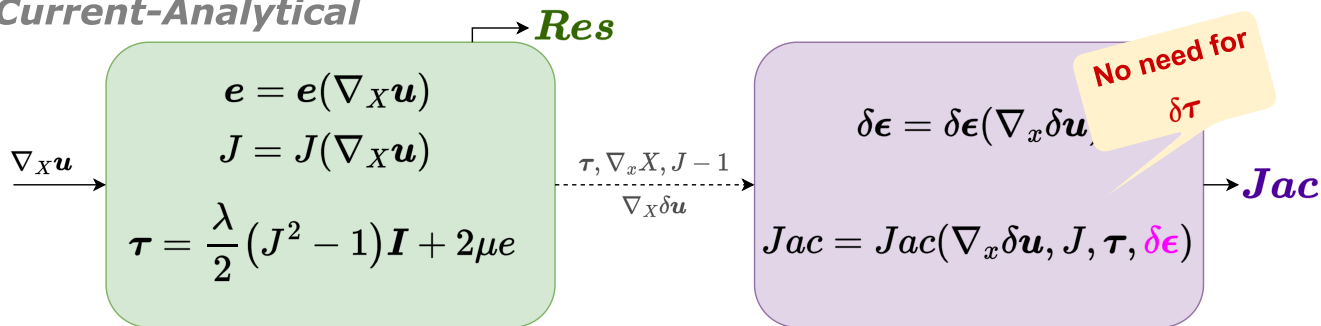
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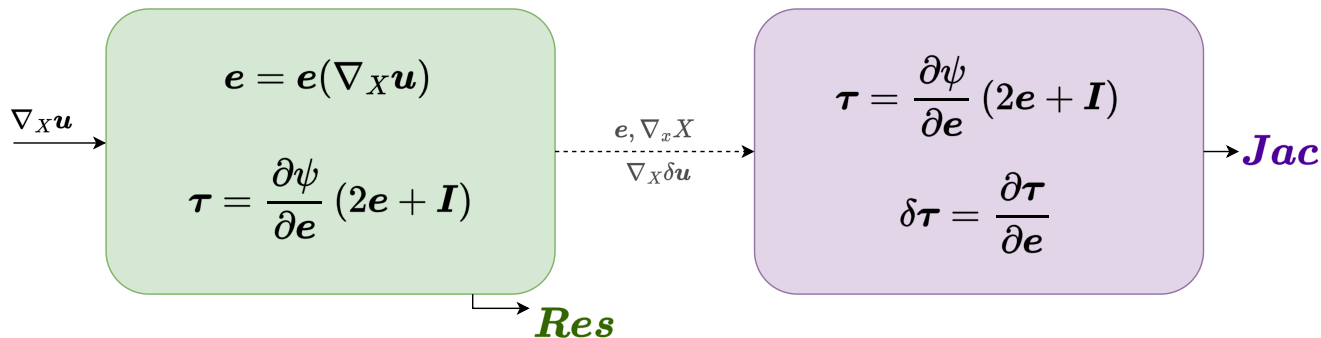
## Current-AD



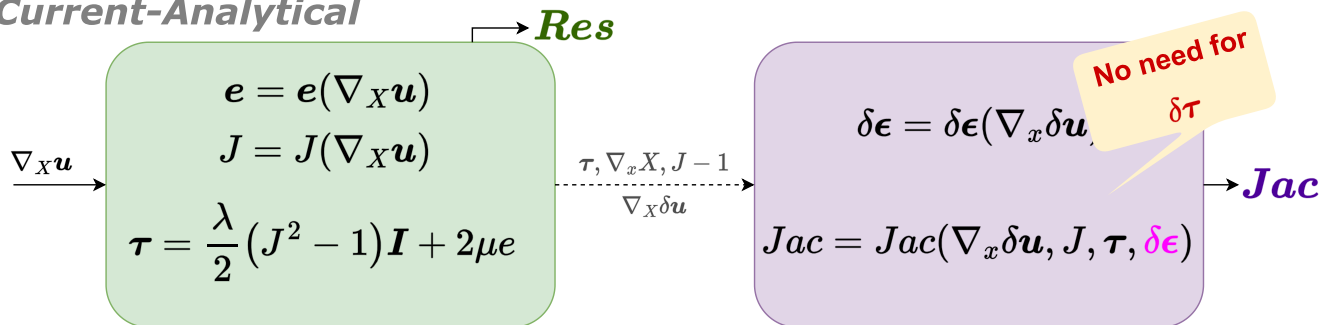
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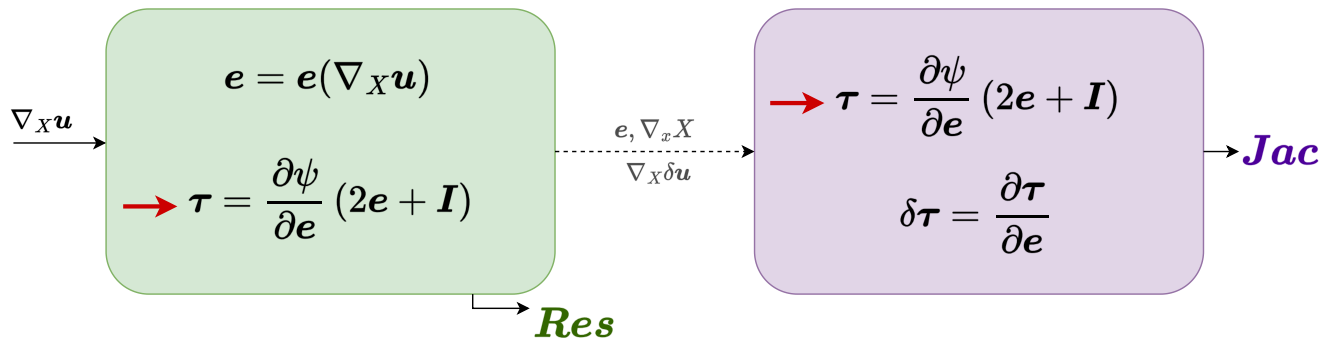
## Current-AD



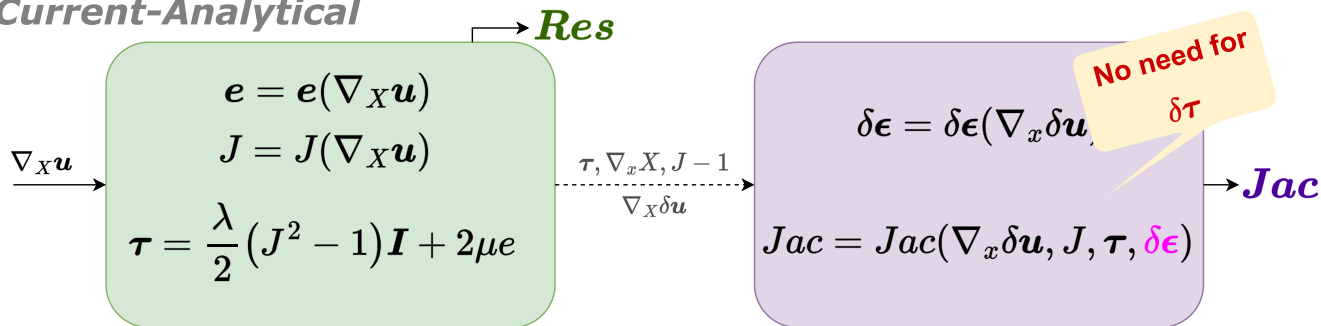
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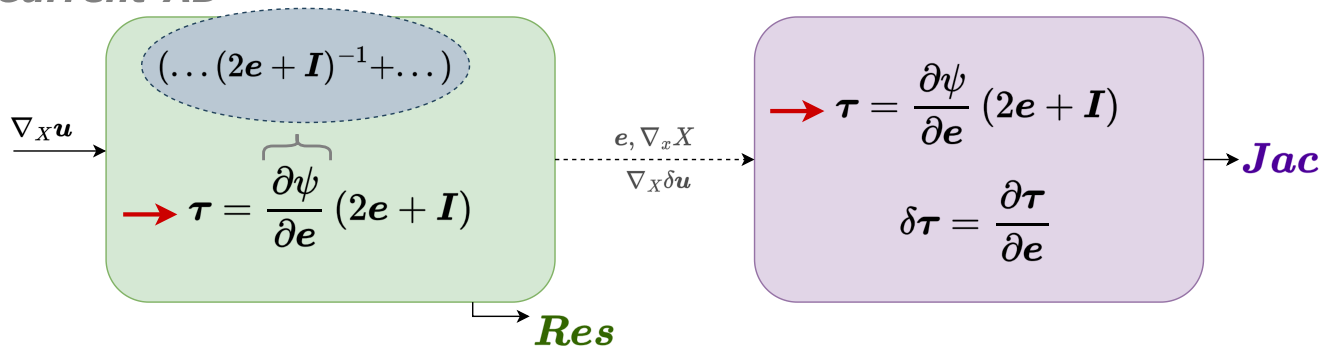
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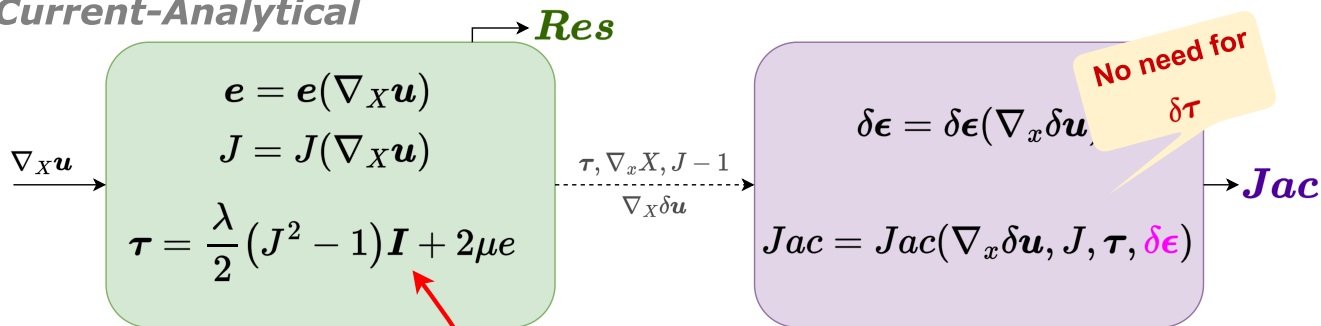
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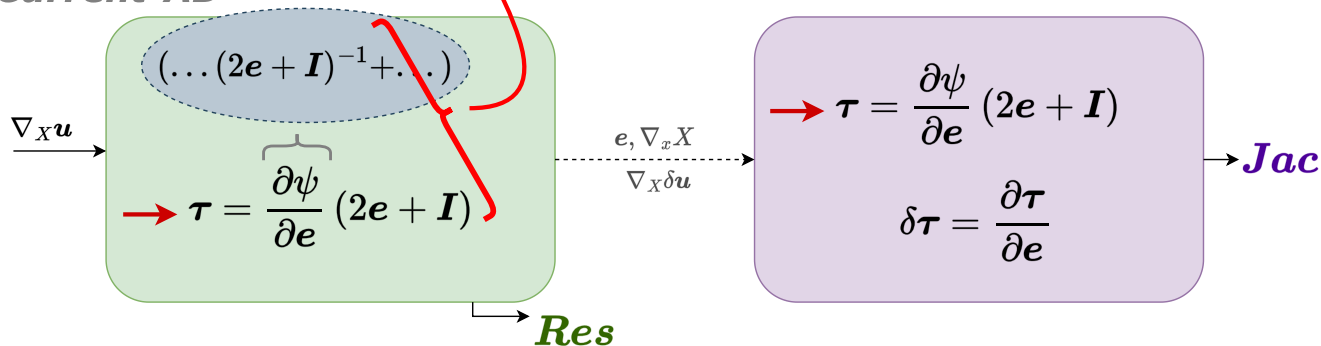
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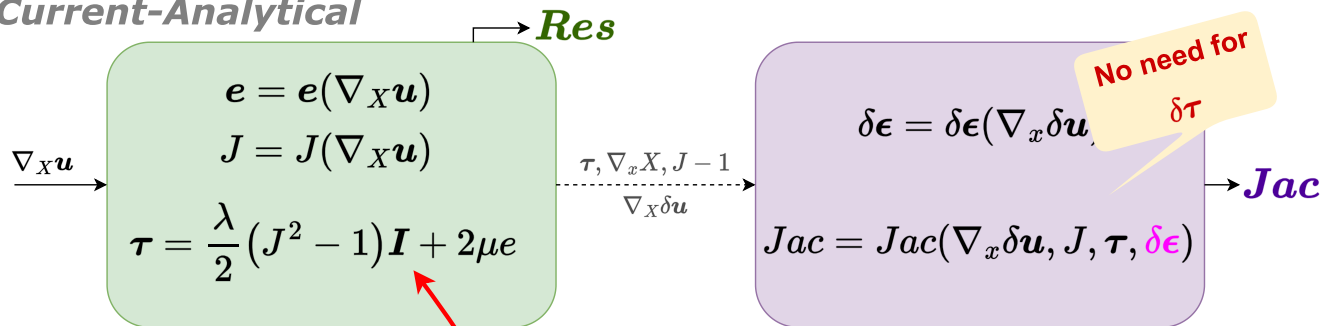
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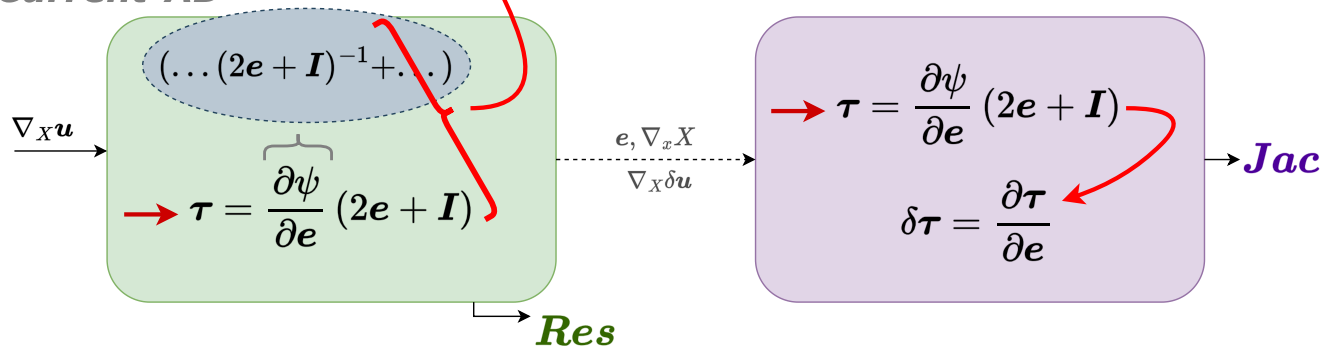
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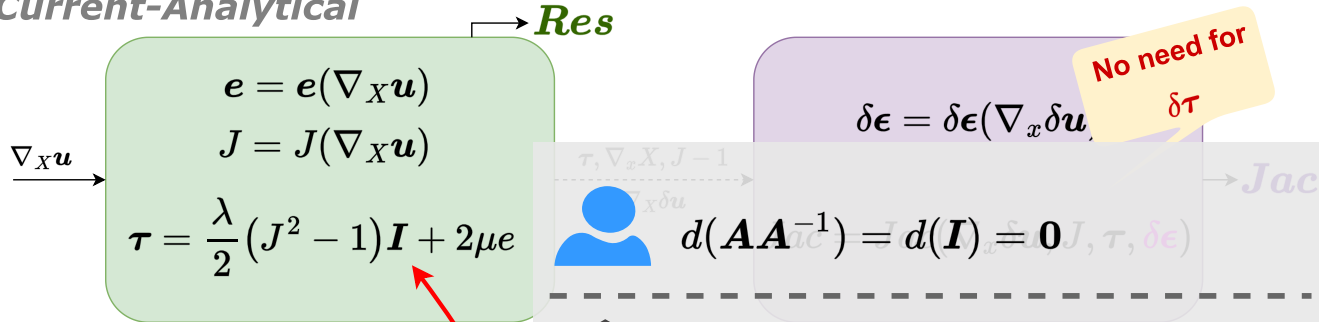


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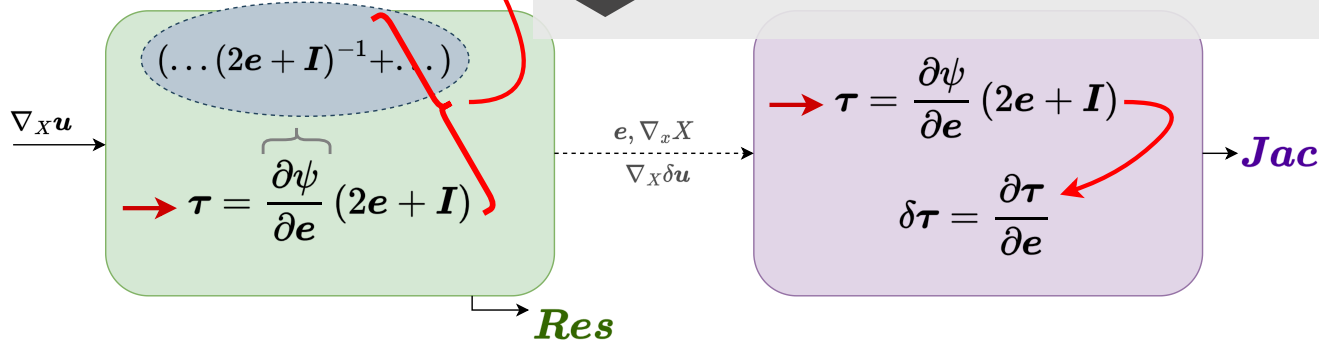




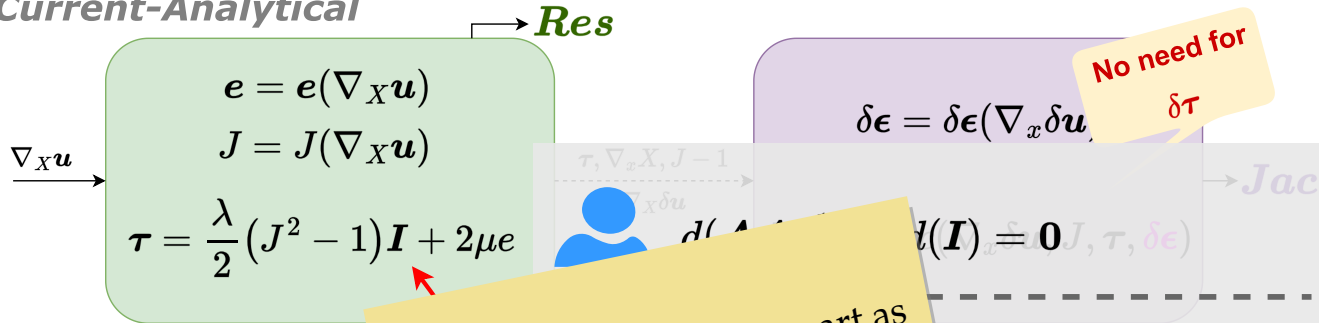
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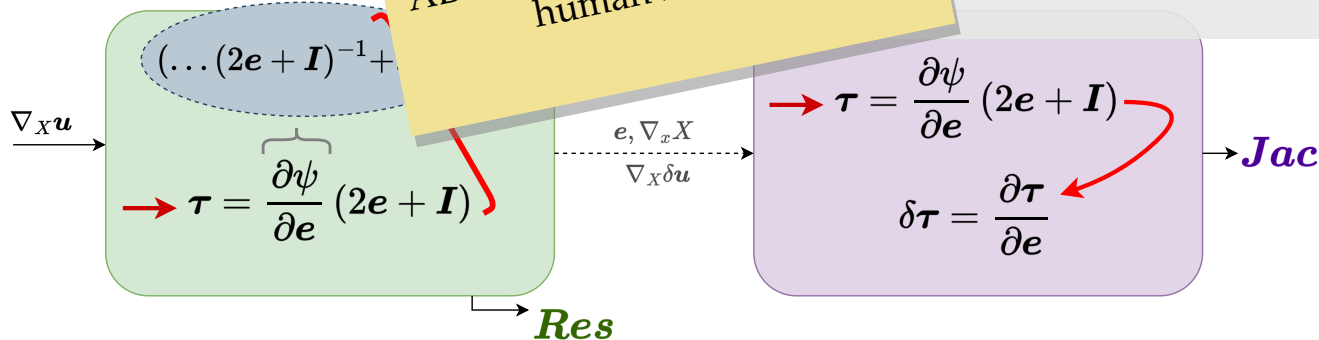
## Current-AD



## Current-Analytical



## Current-AD



# OUTLOOK

## TOWARDS PLASTICITY

$$\text{Input Scalar Functions} = \begin{cases} \psi(\mathbf{E}; \mathbb{I}) & \leftarrow \text{free energy} \\ \phi(\mathbf{S}; \mathbb{I}) & \leftarrow \text{dissipation potential} \\ f(\mathbf{S}; \mathbb{I}) & \leftarrow \text{yield surface} \end{cases}$$

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