

Battery Model Parameter Inference

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1.1 Packages

Here are the packages with brief descriptions (if available):

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ep_bolfi.utility.visualization	
Various helper and plotting functions for common data visualizations	??

2 Hierarchical Index

2.1 Class Hierarchy

This inheritance list is sorted roughly, but not completely, alphabetically:

pybamm.BaseBatteryModel

<code>ep_bolfi.models.electrolyte.Electrolyte</code> <code>pybamm.BaseSubModel</code>	??
<code>ep_bolfi.models.electrolyte.Electrolyte_internal</code>	??
<code>ep_bolfi.models.assess_effective_parameters.Effective_Parameters</code>	??
<code>ep_bolfi.optimization.EP_BOLFI.EP_BOLFI</code> <code>json.JSONEncoder</code>	??
<code>ep_bolfi.optimization.EP_BOLFI.NDArrayEncoder</code>	??
<code>ep_bolfi.utility.fitting_functions.NDArrayEncoder</code>	??
<code>ep_bolfi.utility.dataset_formatting.Measurement</code>	??
<code>ep_bolfi.utility.dataset_formatting.Cycling_Information</code>	??
<code>ep_bolfi.utility.dataset_formatting.Static_Information</code>	??
<code>ep_bolfi.utility.dataset_formatting.Impedance_Measurement</code> object	??
<code>ep_bolfi.utility.fitting_functions.OCV_fit_result</code>	??
<code>ep_bolfi.optimization.EP_BOLFI.Optimizer_State</code>	??
<code>ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator</code> <code>MutableMapping</code>	??
<code>ep_bolfi.utility.preprocessing.SubstitutionDict</code>	??

3 Class Index

3.1 Class List

Here are the classes, structs, unions and interfaces with brief descriptions:

<code>ep_bolfi.utility.dataset_formatting.Cycling_Information</code> Contains basic cycling informations	??
<code>ep_bolfi.models.assess_effective_parameters.Effective_Parameters</code> Calculates, stores and prints effective parameters	??
<code>ep_bolfi.models.electrolyte.Electrolyte</code> Electrolyte model assuming a symmetric Li-metal cell	??
<code>ep_bolfi.models.electrolyte.Electrolyte_internal</code> Defining equations for a symmetric Li cell with electrolyte	??
<code>ep_bolfi.optimization.EP_BOLFI.EP_BOLFI</code> Expectation Propagation and Bayesian Optimization	??
<code>ep_bolfi.utility.dataset_formatting.Impedance_Measurement</code> Contains basic impedance data	??
<code>ep_bolfi.utility.dataset_formatting.Measurement</code> Defines common methods for measurement objects	??

ep_bolfi.optimization.EP_BOLFI.NDArrayEncoder	??
ep_bolfi.utility.fitting_functions.NDArrayEncoder	??
ep_bolfi.utility.fitting_functions.OCV_fit_result Contains OCV fit parameters and related information	??
ep_bolfi.optimization.EP_BOLFI.Optimizer_State Handles the heuristics for the EP-BOLFI operation modes	??
ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator Normalizes sampling to a standard normal distribution	??
ep_bolfi.utility.dataset_formatting.Static_Information Contains additional informations, e.g	??
ep_bolfi.utility.preprocessing.SubstitutionDict A dictionary with some automatic substitutions	??

4 File Index

4.1 File List

Here is a list of all files with brief descriptions:

__init__.py	??
models/__init__.py	??
models/assess_effective_parameters.py	??
models/electrolyte.py	??
models/solversetup.py	??
models/standard_parameters.py	??
optimization/__init__.py	??
optimization/EP_BOLFI.py	??
utility/__init__.py	??
utility/dataset_formatting.py	??
utility/fitting_functions.py	??
utility/preprocessing.py	??
utility/visualization.py	??

5 Namespace Documentation

5.1 ep_bolfi Namespace Reference

Namespaces

- [models](#)
- [optimization](#)
- [utility](#)

5.2 `ep_bolfi`.`EP_BOLFI`.`EP_BOLFI` Namespace Reference

This file contains functions to perform Expectation Propagation on simulator models using BOLFI (Bayesian Optimization).

5.2.1 Detailed Description

This file contains functions to perform Expectation Propagation on simulator models using BOLFI (Bayesian Optimization).

5.3 `ep_bolfi.models` Namespace Reference

Namespaces

- [`assess_effective_parameters`](#)
Evaluates a parameter set for, e.g., overpotential and capacity.
- [`electrolyte`](#)
Contains a PyBaMM-compatible electrolyte model.
- [`solversetup`](#)
This file eases the setup and simulation of PyBaMM battery models.
- [`standard_parameters`](#)
Comprehensive list of parameters of every model.

5.4 `ep_bolfi.models.assess_effective_parameters` Namespace Reference

Evaluates a parameter set for, e.g., overpotential and capacity.

Classes

- class [`Effective_Parameters`](#)
Calculates, stores and prints effective parameters.

5.4.1 Detailed Description

Evaluates a parameter set for, e.g., overpotential and capacity.

5.5 `ep_bolfi.models.electrolyte` Namespace Reference

Contains a PyBaMM-compatible electrolyte model.

Classes

- class [`Electrolyte_internal`](#)
Defining equations for a symmetric Li cell with electrolyte.
- class [`Electrolyte`](#)
[Electrolyte](#) model assuming a symmetric Li-metal cell.

5.5.1 Detailed Description

Contains a PyBaMM-compatible electrolyte model.

5.6 ep_bolfi.models.solversetup Namespace Reference

This file eases the setup and simulation of PyBaMM battery models.

Functions

- def [solver_setup](#) (model, parameters, submesh_types, var_pts, spatial_methods, geometry=None, reltol=1e-6, abstol=1e-6, root_tol=1e-3, dt_max=None, free_parameters=[], verbose=False, logging_file=None)
Processes the model and returns a runnable solver.
- def [simulation_setup](#) (model, operation_input, parameters, submesh_types, var_pts, spatial_methods, geometry=None, reltol=1e-6, abstol=1e-6, root_tol=1e-3, dt_max=None, free_parameters=[], verbose=False, logging_file=None)
Processes the model and returns a runnable solver.
- def [auto_var_pts](#) (x_n, x_s, x_p, r_n, r_p, y=1, z=1)
Utility function for setting the discretization density.
- def [spectral_mesh_pts_and_method](#) (order_s_n, order_s_p, order_e, volumes_e_n=1, volumes_e_s=1, volumes_e_p=1, halfcell=False)
Utility function for default mesh and spatial methods.

5.6.1 Detailed Description

This file eases the setup and simulation of PyBaMM battery models.

5.6.2 Function Documentation

5.6.2.1 auto_var_pts() def ep_bolfi.models.solversetup.auto_var_pts (
 x_n,
 x_s,
 x_p,
 r_n,
 r_p,
 y = 1,
 z = 1)

Utility function for setting the discretization density.

Parameters

x_n	The number of voxels for the electrolyte in the anode.
x_s	The number of voxels for the electrolyte in the separator.

Parameters

$x \leftrightarrow$ _p	The number of voxels for the electrolyte in the cathode.
$r \leftrightarrow$ _n	The number of voxels for each anode representative particle.
$r \leftrightarrow$ _p	The number of voxels for each cathode representative particle.
y	Used by PyBaMM for spatially resolved current collectors. Don't change the default (1) unless the model supports it.
z	Used by PyBaMM for spatially resolved current collectors. Don't change the default (1) unless the model supports it.

Returns

A discretization dictionary that can be used with PyBaMM models.

5.6.2.2 simulation_setup() `def ep_bolfi.models.solversetup.simulation_setup (`
 model,
 operation_input,
 parameters,
 submesh_types,
 var_pts,
 spatial_methods,
 geometry = None,
 reltol = 1e-6,
 abstol = 1e-6,
 root_tol = 1e-3,
 dt_max = None,
 free_parameters = [],
 verbose = False,
 logging_file = None)

Processes the model and returns a runnable solver.

In contrast to [solversetup.solver_setup](#), this allows for a more verbose description of the operating conditions and should be preferred.

Parameters

<i>model</i>	A PyBaMM model. Use one of the models in this folder.
<i>operation_input</i>	A list of strings which describe the operating conditions. These exactly match the PyBaMM usage for <code>pybamm.Experiment</code> . Examples: "Hold at 4 V for 60 s", "Discharge at 1 A for 30 s", "Rest for 1800 s", "Charge at 1 C for 30 s".
<i>parameters</i>	The parameters that the model requires as a dictionary. Please refer to models.standard_parameters for the names or adapt one of the examples in <code>parameters.models</code> .
<i>submesh_types</i>	A dictionary of the meshes to be used. The keys have to match the geometry names in the model. Use <code>#spectral_mesh_and_method</code> as reference or a shortcut.
<i>var_pts</i>	A dictionary giving the number of discretization volumes. Since the keys have to be special variables determined by PyBaMM, use auto_var_pts as a shortcut.

Parameters

<i>spatial_methods</i>	A dictionary of the spatial methods to be used. The keys have to match the geometry names in the model. Use #spectral_mesh_and_method as reference or a shortcut.
<i>geometry</i>	The geometry of the model in dictionary form. Usually, model.default_geometry is sufficient, which is the default.
<i>reltol</i>	The relative tolerance that the Casadi solver should use. Default is 1e-6.
<i>abstol</i>	The absolute tolerance that the Casadi solver should use. Default is 1e-6.
<i>root_tol</i>	The tolerance for rootfinding that the Casadi solver should use. Default is 1e-3.
<i>dt_max</i>	The maximum timestep size for the Casadi solver in seconds. Default is chosen by PyBaMM.
<i>free_parameters</i>	A list of parameter names that shall be input later. They may be given to the solve() function of the returned Simulation object as a dictionary to the keyword parameter "inputs" with the names as keys and the values of the parameters as values. DO NOT USE GEOMETRICAL PARAMETERS, THEY WILL CRASH THE MESH. Instead, just use this function with a complete set of parameters where the relevant parameters are changed.
<i>verbose</i>	The default (False) sets the PyBaMM flag to only show warnings. True will show the details of preprocessing and the runtime of the solver. This applies globally, so don't set this to True if running simulations in parallel.
<i>logging_file</i>	Optional name of a file to store the logs in.

Returns

A 2-tuple of a pybamm.Simulation.solve call that runs the simulation when called, and the proper callback for the logging file if specified (else None). Please use solve(check_model=False) if it doesn't work properly with redundant model checks in place.

5.6.2.3 solver_setup() def ep_bolfi.models.solversetup.solver_setup (
model,
parameters,
submesh_types,
var_pts,
spatial_methods,
geometry = None,
reltol = 1e-6,
abstol = 1e-6,
root_tol = 1e-3,
dt_max = None,
free_parameters = [],
verbose = False,
logging_file = None)

Processes the model and returns a runnable solver.

Parameters

<i>model</i>	A PyBaMM model. Use one of the models in this folder.
<i>parameters</i>	The parameters that the model requires as a dictionary. Please refer to models.standard_parameters for the names or adapt one of the examples in parameters.models.

Parameters

<i>submesh_types</i>	A dictionary of the meshes to be used. The keys have to match the geometry names in the model. Use #spectral_mesh_and_method as reference or a shortcut.
<i>var_pts</i>	A dictionary giving the number of discretization volumes. Since the keys have to be special variables determined by PyBaMM, use auto_var_pts as a shortcut.
<i>spatial_methods</i>	A dictionary of the spatial methods to be used. The keys have to match the geometry names in the model. Use #spectral_mesh_and_method as reference or a shortcut.
<i>geometry</i>	The geometry of the model in dictionary form. Usually, model.default_geometry is sufficient, which is the default.
<i>reltol</i>	The relative tolerance that the Casadi solver shall use. Default is 1e-6.
<i>abstol</i>	The absolute tolerance that the Casadi solver shall use. Default is 1e-6.
<i>root_tol</i>	The tolerance for rootfinding that the Casadi solver shall use. Default is 1e-3.
<i>dt_max</i>	The maximum timestep size for the Casadi solver in seconds. Default is chosen by PyBaMM.
<i>free_parameters</i>	A list of parameter names that shall be input later. They may be given to the returned lambda function as a dictionary with the names as keys and the values of the parameters as values. DO NOT USE GEOMETRICAL PARAMETERS, THEY WILL CRASH THE MESH. Instead, just use this function with a complete set of parameters where the relevant parameters are changed.
<i>verbose</i>	The default (False) sets the PyBaMM flag to only show warnings. True will show the details of preprocessing and the runtime of the solver. This applies globally, so don't set this to True if running simulations in parallel.
<i>logging_file</i>	Optional name of a file to store the logs in.

Returns

A lambda function that takes a numpy.array of timepoints to evaluate and runs the Casadi solver for those. Optionally takes a dictionary of parameters as specified by "free_parameters".

5.6.2.4 spectral_mesh_pts_and_method() `def ep_bolfi.models.solversetup.spectral_mesh_pts_and_method (`
`order_s_n,`
`order_s_p,`
`order_e,`
`volumes_e_n = 1,`
`volumes_e_s = 1,`
`volumes_e_p = 1,`
`halfcell = False)`

Utility function for default mesh and spatial methods.

Only returns Spectral Volume mesh and spatial methods.

Parameters

<i>order_s_n</i>	The order of the anode particles Spectral Volumes.
<i>order_s_p</i>	The order of the anode particles Spectral Volumes.
<i>order_e</i>	The order of the anode, separator and cathode electrolyte Spectral Volumes. These have to be the same, since the corresponding meshes get concatenated.

Parameters

<code>volumes_e↔ _n</code>	The # of Spectral Volumes to use for the anode electrolyte. This is useful to have different resolutions for each part.
<code>volumes_e↔ _s</code>	The # of Spectral Volumes to use for the separator electrolyte.
<code>volumes_e↔ _p</code>	The # of Spectral Volumes to use for the cathode electrolyte.
<code>halfcell</code>	Default is False, which sets up the mesh and spatial methods for a full-cell setup. Set it to True for a half-cell setup.

Returns

A (submesh_types, spatial_methods) tuple for PyBaMM usage.

5.7 `ep_bolfi.models.standard_parameters` Namespace Reference

Comprehensive list of parameters of every model.

Functions

- def `SOC_n_dim_init` (x)
Initial SOC of the anode.
- def `SOC_n_init` (x)
Non-dimensionalized initial SOC of the anode.
- def `SOC_p_dim_init` (x)
Initial SOC of the cathode.
- def `SOC_p_init` (x)
Non-dimensionalized initial SOC of the cathode.
- def `D_e_dim` (c_e_dim, T_dim)
Electrolyte diffusivity.
- def `D_e` (c_e, T)
Non-dimensionalized electrolyte diffusivity.
- def `κ_e_dim` (c_e_dim, T_dim)
Electrolyte conductivity.
- def `κ_e` (c_e, T)
Non-dimensionalized electrolyte conductivity.
- def `t_plus_dim` (c_e_dim)
Transference number.
- def `t_plus` (c_e)
Non-dimensionalized (referring to the input) transference number.
- def `one_plus_dlnf_dlnc_dim` (c_e_dim)
Thermodynamic factor.
- def `one_plus_dlnf_dlnc` (c_e)
Non-dimensionalized (referring to the input) thermodynamic factor.
- def `D_n_dim` (SOC_n, T_dim)
Anode diffusivity.
- def `D_n` (SOC_n, T)
Non-dimensionalized anode diffusivity.

- def D_{p_dim} (SOC_p , T_dim)
Cathode diffusivity.
- def D_p (SOC_p , T)
Non-dimensionalized cathode diffusivity.
- def $i_{sa_0_dim}$ (c_a_dim , $SOC_{n_surf_dim}$, c_{n_max} , T_dim)
Anode exchange current density.
- def i_{sa_0} (c_a , SOC_{n_surf} , c_{n_max} , T)
Non-dimensionalized anode exchange current density.
- def $d_c_a_i_{sa_0_dim}$ (c_a_dim , $SOC_{n_surf_dim}$, c_{n_max} , T_dim)
 ∂ anode exchange current density / ∂ electrolyte concentration.
- def $d_c_a_i_{sa_0}$ (c_a , SOC_{n_surf} , c_{n_max} , T)
The non-dimensionalized version of the prior variable.
- def $i_{sep_0_dim}$ (c_{ep_dim} , $SOC_{p_surf_dim}$, c_{p_max} , T_dim)
Cathode exchange current density.
- def i_{sep_0} (c_{ep} , SOC_{p_surf} , c_{p_max} , T)
Non-dimensionalized cathode exchange current density.
- def $d_c_{ep_i_{sep_0_dim}}$ (c_{ep_dim} , $SOC_{p_surf_dim}$, c_{p_max} , T_dim)
 ∂ cathode exchange current density / ∂ electrolyte concentration.
- def $d_c_{ep_i_{sep_0}}$ (c_{ep} , SOC_{p_surf} , c_{p_max} , T)
The non-dimensionalized version of the prior variable.
- def $dOCV_{n_dT_dim}$ (SOC_n)
 ∂ anode OCV / ∂ temperature.
- def $dOCV_{n_dT}$ (SOC_n)
Non-dimensionalized ∂ anode OCV / ∂ temperature.
- def $dOCV_{n_dT_dSOC_n_dim}$ (SOC_n)
 $(\partial$ anode OCV / ∂ temperature) / ∂ anode SOC.
- def $dOCV_{n_dT_dSOC_n}$ (SOC_n)
Non-dimensionalized $(\partial$ anode OCV / ∂ temperature) / ∂ anode SOC.
- def $dOCV_{p_dT_dim}$ (SOC_p)
 ∂ cathode OCV / ∂ temperature.
- def $dOCV_{p_dT}$ (SOC_p)
Non-dimensionalized ∂ cathode OCV / ∂ temperature.
- def $dOCV_{p_dT_dSOC_p_dim}$ (SOC_p)
 $(\partial$ cathode OCV / ∂ temperature) / ∂ cathode SOC.
- def $dOCV_{p_dT_dSOC_p}$ (SOC_p)
Non-dimensionalized $(\partial$ cathode OCV / ∂ temperature) / ∂ cathode SOC.
- def OCV_{n_dim} (SOC_n , T_dim)
Anode OCV.
- def OCV_n (SOC_n , T)
Non-dimensionalized anode OCV.
- def $dOCV_{n_dim_dSOC_n}$ (SOC_n , T_dim)
 ∂ anode OCV / ∂ anode SOC.
- def $dOCV_{n_dSOC_n}$ (SOC_n , T)
Non-dimensionalized ∂ anode OCV / ∂ anode SOC.
- def OCV_{p_dim} (SOC_p , T_dim)
Cathode OCV.
- def OCV_p (SOC_p , T)
Non-dimensionalized cathode OCV.
- def $dOCV_{p_dim_dSOC_p}$ (SOC_p , T_dim)
 ∂ cathode OCV / ∂ cathode SOC.
- def $dOCV_{p_dSOC_p}$ (SOC_p , T)
Non-dimensionalized ∂ cathode OCV / ∂ cathode SOC.

Variables

- R = Scalar(constants.R)
- F = Scalar(constants.physical_constants["Faraday constant"][0])
- k_B = constants.physical_constants["Boltzmann constant"][0]
- q_e = constants.physical_constants["electron volt"][0]
- T_{ref} = Parameter("Reference temperature [K]")
- T_{init} = Parameter("Initial temperature [K]")
- $thermal_voltage = R * T_{ref} / F$
- ΔT = Scalar(1)
- L_{n_dim} = Parameter("Negative electrode thickness [m]")
- L_{s_dim} = Parameter("Separator thickness [m]")
- L_{p_dim} = Parameter("Positive electrode thickness [m]")
- $L_{dim} = L_{n_dim} + L_{s_dim} + L_{p_dim}$
- A = Parameter("Current collector perpendicular area [m2]")
- V = Parameter("Cell volume [m3]")
- $L_x = L_{dim}$
- L_y = Parameter("Electrode width [m]")
- L_z = Parameter("Electrode height [m]")
- C = Parameter("Typical current [A]")
- U_l = pybamm.Parameter("Lower voltage cut-off [V]")
- U_u = pybamm.Parameter("Upper voltage cut-off [V]")
- c_{e_typ} = pybamm.Parameter("Typical electrolyte concentration [mol.m-3]")
- c_n = pybamm.Parameter("Maximum concentration in negative electrode [mol.m-3]")
- c_p = pybamm.Parameter("Maximum concentration in positive electrode [mol.m-3]")
- σ_{n_dim} = pybamm.Parameter("Negative electrode conductivity [S.m-1]")
- σ_{p_dim} = pybamm.Parameter("Positive electrode conductivity [S.m-1]")
- a_{n_dim} = Parameter("Negative electrode surface area to volume ratio [m-1]")
- a_{p_dim} = Parameter("Positive electrode surface area to volume ratio [m-1]")
- R_n = Parameter("Negative particle radius [m]")
- R_p = Parameter("Positive particle radius [m]")
- α_{nn} = Parameter("Negative electrode anodic charge-transfer coefficient")
- α_{pn} = Parameter("Negative electrode cathodic charge-transfer coefficient")
- α_{np} = Parameter("Positive electrode anodic charge-transfer coefficient")
- α_{pp} = Parameter("Positive electrode cathodic charge-transfer coefficient")
- β_n_scalar
- β_{es_scalar} = Parameter("Separator Bruggeman coefficient (electrolyte)")
- β_{ep_scalar}
- β_n = pybamm.PrimaryBroadcast(β_n_scalar , "negative electrode")
- β_{es} = pybamm.PrimaryBroadcast(β_{es_scalar} , "separator")
- β_{ep} = pybamm.PrimaryBroadcast(β_{ep_scalar} , "positive electrode")
- β_{sn_scalar} = Parameter("Negative electrode Bruggeman coefficient (electrode)")
- β_{ss_scalar} = Parameter("Separator Bruggeman coefficient (electrode)")
- β_{sp_scalar} = Parameter("Positive electrode Bruggeman coefficient (electrode)")
- β_e = pybamm.Concatenation(β_n , β_{es} , β_{ep})
- ϵ_n_scalar = Parameter("Negative electrode porosity")
- ϵ_s_scalar = Parameter("Separator porosity")
- ϵ_p_scalar = Parameter("Positive electrode porosity")
- ϵ_n = pybamm.PrimaryBroadcast(ϵ_n_scalar , "negative electrode")
- ϵ_s = pybamm.PrimaryBroadcast(ϵ_s_scalar , "separator")
- ϵ_p = pybamm.PrimaryBroadcast(ϵ_p_scalar , "positive electrode")
- ϵ = pybamm.Concatenation(ϵ_n , ϵ_s , ϵ_p)
- $\epsilon^\beta = \epsilon ** \beta_e$
- z_n = Parameter("Negative electrode electrons in reaction")
- z_p = Parameter("Positive electrode electrons in reaction")

- `c_e_dim_init` = Parameter("Initial concentration in electrolyte [mol.m-3]")
- `c_e_init` = `c_e_dim_init` / `c_e_typ`
- `def D_e_typ` = `D_e_dim`(`c_e_typ`, `T_ref`)
- `def κ_e_typ` = `κ_e_dim`(`c_e_typ`, `T_ref`)
- `tuple κ_e_hat` = (`R * T_ref / F`) / (`C / A * L_dim / κ_e_typ`)
- `def D_n_typ` = `D_n_dim`(`Scalar(0.5)`, `T_ref`)
- `def D_p_typ` = `D_p_dim`(`Scalar(0.5)`, `T_ref`)
- `def i_sn_0_ref` = `i_sn_0_dim`(`c_e_typ`, `0.5 * c_n`, `c_n`, `T_ref`)
- `def i_sep_0_ref` = `i_sep_0_dim`(`c_e_typ`, `0.5 * c_p`, `c_p`, `T_ref`)
- `def OCV_n_ref` = `OCV_n_dim`(`SOC_n_init(0)`, `T_ref`)
- `def OCV_p_ref` = `OCV_p_dim`(`SOC_p_init(1)`, `T_ref`)
- `a_n` = `a_n_dim` * `R_n`
- `a_p` = `a_p_dim` * `R_p`
- `tuple Q` = (`1 - ε_p_scalar`) * `L_p_dim` * `c_p` * `z_p` * `F` * `A`
- `τd` = `F * c_p * L_dim` / (`C / A`)
- `int τ_e` = `L_dim**2 / D_e_typ`
- `int τ_n` = `R_n**2 / D_n_typ`
- `int τ_p` = `R_p**2 / D_p_typ`
- `τrn` = `F * c_n` / (`i_sn_0_ref * a_n_dim`)
- `τrp` = `F * c_p` / (`i_sep_0_ref * a_p_dim`)
- `timescale` = `τd`
- `int C_e` = `τ_e / τd`
- `int C_n` = `τ_n / τd`
- `int C_p` = `τ_p / τd`
- `Crn` = `τrn / τd`
- `Crp` = `τrp / τd`
- `γ_e` = `c_e_typ` / `c_p`
- `γ_n` = `c_n` / `c_p`
- `γ_p` = `c_p` / `c_p`
- `Ln` = `L_n_dim` / `L_dim`
- `Ls` = `L_s_dim` / `L_dim`
- `Lp` = `L_p_dim` / `L_dim`
- `Le`
- `tuple σn` = (`thermal_voltage / (C / A * L_dim)`) * `σn_dim`
- `tuple σp` = (`thermal_voltage / (C / A * L_dim)`) * `σp_dim`
- `Iextern_dim`
- `Iextern` = `Iextern_dim / C`
- `nelectrodes_parallel`
- `ncells` = Parameter("Number of cells connected in series to make a battery")
- `Ityp` = `C`
- `Acc` = `A`
- `current_with_time` = `Iextern`
- `dimensional_current_with_time` = `Iextern_dim`
- `dimensional_current_density_with_time` = `Iextern_dim / A`
- `voltage_low_cut` = `Ul`
- `voltage_high_cut` = `Uu`
- `capacity` = Parameter("Nominal cell capacity [A.h]")

5.7.1 Detailed Description

Comprehensive list of parameters of every model.

SI units are assumed unless stated otherwise. When imported, this package turns into the list of variables contained in here. The "syntactic sugar" with the lower and upper indexed letters is treated by Python to be the same as their normal counterparts. E.g., "`an`" and "`an`" refer to the exact same variable. Greek letters aren't converted, unless they are also indexed, in which case the same as before applies: "`εβ`" is the same as "`εβ`".

5.7.2 Function Documentation

5.7.2.1 d_c_ai_{sa}0() def ep_bolfi.models.standard_parameters.d_c_ai_{sa}0 (
 c_a,
 SOC_n_surf,
 c_n_max,
 T)

The non-dimensionalized version of the prior variable.

5.7.2.2 d_c_ai_{sa}0_dim() def ep_bolfi.models.standard_parameters.d_c_ai_{sa}0_dim (
 c_a_dim,
 SOC_n_surf_dim,
 c_n_max,
 T_dim)

∂ anode exchange current density / ∂ electrolyte concentration.

5.7.2.3 d_c_pi_{sep}0() def ep_bolfi.models.standard_parameters.d_c_pi_{sep}0 (
 c_{ep},
 SOC_p_surf,
 c_p_max,
 T)

The non-dimensionalized version of the prior variable.

5.7.2.4 d_c_pi_{sep}0_dim() def ep_bolfi.models.standard_parameters.d_c_pi_{sep}0_dim (
 c_{ep}_dim,
 SOC_p_surf_dim,
 c_p_max,
 T_dim)

∂ cathode exchange current density / ∂ electrolyte concentration.

5.7.2.5 dOCV_n_dim_dSOC_n() def ep_bolfi.models.standard_parameters.dOCV_n_dim_dSOC_n (
 SOC_n,
 T_dim)

∂ anode OCV / ∂ anode SOC.

5.7.2.6 dOCV_ndSOC_n() def ep_bolfi.models.standard_parameters.dOCV_ndSOC_n (
 SOC_n,
 T)

Non-dimensionalized ∂ anode OCV / ∂ anode SOC.

5.7.2.7 dOCV_ndT() def ep_bolfi.models.standard_parameters.dOCV_ndT (
 SOC_n)

Non-dimensionalized ∂ anode OCV / ∂ temperature.

5.7.2.8 dOCV_ndT_{dim}() def ep_bolfi.models.standard_parameters.dOCV_ndT_{dim} (
 SOC_n)

∂ anode OCV / ∂ temperature.

5.7.2.9 dOCV_ndT_{dim}dSOC_n() def ep_bolfi.models.standard_parameters.dOCV_ndT_{dim}dSOC_n (
 SOC_n)

Non-dimensionalized (∂ anode OCV / ∂ temperature) / ∂ anode SOC.

5.7.2.10 dOCV_ndT_{dim}dSOC_ndim() def ep_bolfi.models.standard_parameters.dOCV_ndT_{dim}dSOC_ndim (
 SOC_n)

(∂ anode OCV / ∂ temperature) / ∂ anode SOC.

5.7.2.11 dOCV_pdim_{dim}dSOC_p() def ep_bolfi.models.standard_parameters.dOCV_pdim_{dim}dSOC_p (
 SOC_p,
 T_{dim})

∂ cathode OCV / ∂ cathode SOC.

5.7.2.12 dOCV_pdSOC_p() def ep_bolfi.models.standard_parameters.dOCV_pdSOC_p (
 SOC_p,
 T)

Non-dimensionalized ∂ cathode OCV / ∂ cathode SOC.

5.7.2.13 dOCV_p_dT() def ep_bolfi.models.standard_parameters.dOCV_p_dT (
 SOC_p)

Non-dimensionalized ∂ cathode OCV / ∂ temperature.

5.7.2.14 dOCV_p_dT_dim() def ep_bolfi.models.standard_parameters.dOCV_p_dT_dim (
 SOC_p)

∂ cathode OCV / ∂ temperature.

5.7.2.15 dOCV_p_dT_dSOC_p() def ep_bolfi.models.standard_parameters.dOCV_p_dT_dSOC_p (
 SOC_p)

Non-dimensionalized (∂ cathode OCV / ∂ temperature) / ∂ cathode SOC.

5.7.2.16 dOCV_p_dT_dSOC_p_dim() def ep_bolfi.models.standard_parameters.dOCV_p_dT_dSOC_p_dim (
 SOC_p)

(∂ cathode OCV / ∂ temperature) / ∂ cathode SOC.

5.7.2.17 D_e() def ep_bolfi.models.standard_parameters.D_e (
 c_e,
 T)

Non-dimensionalized electrolyte diffusivity.

5.7.2.18 D_e_dim() def ep_bolfi.models.standard_parameters.D_e_dim (
 c_{e_dim},
 T_dim)

Electrolyte diffusivity.

5.7.2.19 D_n() def ep_bolfi.models.standard_parameters.D_n (
 SOC_n,
 T)

Non-dimensionalized anode diffusivity.

5.7.2.20 $D_n_dim()$ `def ep_bolfi.models.standard_parameters.Dn_dim (`
 `SOCn,`
 `T_dim)`

Anode diffusivity.

5.7.2.21 $D_p()$ `def ep_bolfi.models.standard_parameters.Dp (`
 `SOCp,`
 `T)`

Non-dimensionalized cathode diffusivity.

5.7.2.22 $D_p_dim()$ `def ep_bolfi.models.standard_parameters.Dp_dim (`
 `SOCp,`
 `T_dim)`

Cathode diffusivity.

5.7.2.23 $i_{se_0}()$ `def ep_bolfi.models.standard_parameters.i_se_0 (`
 `ca,`
 `SOCn_surf,`
 `cn_max,`
 `T)`

Non-dimensionalized anode exchange current density.

5.7.2.24 $i_{se_0_dim}()$ `def ep_bolfi.models.standard_parameters.i_se_0_dim (`
 `ca_dim,`
 `SOCn_surf_dim,`
 `cn_max,`
 `T_dim)`

Anode exchange current density.

5.7.2.25 $i_{sep_0}()$ `def ep_bolfi.models.standard_parameters.i_sep_0 (`
 `cep,`
 `SOCp_surf,`
 `cp_max,`
 `T)`

Non-dimensionalized cathode exchange current density.

5.7.2.26 i_sep_0_dim() def ep_bolfi.models.standard_parameters.i_sep_0_dim (
 c_{ep_dim},
 SOC_{p_surf_dim},
 c_{p_max},
 T_dim)

Cathode exchange current density.

5.7.2.27 OCV_n() def ep_bolfi.models.standard_parameters.OCV_n (
 SOC_n,
 T)

Non-dimensionalized anode OCV.

5.7.2.28 OCV_{n_dim}() def ep_bolfi.models.standard_parameters.OCV_{n_dim} (
 SOC_n,
 T_dim)

Anode OCV.

5.7.2.29 OCV_p() def ep_bolfi.models.standard_parameters.OCV_p (
 SOC_p,
 T)

Non-dimensionalized cathode OCV.

5.7.2.30 OCV_{p_dim}() def ep_bolfi.models.standard_parameters.OCV_{p_dim} (
 SOC_p,
 T_dim)

Cathode OCV.

5.7.2.31 one_plus_dlnf_dlnc() def ep_bolfi.models.standard_parameters.one_plus_dlnf_dlnc (
 c_e)

Non-dimensionalized (referring to the input) thermodynamic factor.

5.7.2.32 one_plus_dlnf_dlnc_dim() def ep_bolfi.models.standard_parameters.one_plus_dlnf_dlnc_dim (
 c_e dim)

Thermodynamic factor.

5.7.2.33 SOC_n_dim_init() def ep_bolfi.models.standard_parameters.SOC_n_dim_init (
 x)

Initial SOC of the anode.

5.7.2.34 SOC_n_init() def ep_bolfi.models.standard_parameters.SOC_n_init (
 x)

Non-dimensionalized initial SOC of the anode.

5.7.2.35 SOC_p_dim_init() def ep_bolfi.models.standard_parameters.SOC_p_dim_init (
 x)

Initial SOC of the cathode.

5.7.2.36 SOC_p_init() def ep_bolfi.models.standard_parameters.SOC_p_init (
 x)

Non-dimensionalized initial SOC of the cathode.

5.7.2.37 t_plus() def ep_bolfi.models.standard_parameters.t_plus (
 c_e)

Non-dimensionalized (referring to the input) transference number.

5.7.2.38 t_plus_dim() def ep_bolfi.models.standard_parameters.t_plus_dim (
 c_e dim)

Transference number.

5.7.2.39 $\kappa_e()$ def ep_bolfi.models.standard_parameters. κ_e (
 c_e ,
 T)

Non-dimensionalized electrolyte conductivity.

5.7.2.40 $\kappa_{e_dim}()$ def ep_bolfi.models.standard_parameters. κ_{e_dim} (
 c_{e_dim} ,
 T_{dim})

Electrolyte conductivity.

5.7.3 Variable Documentation

5.7.3.1 A ep_bolfi.models.standard_parameters. A = Parameter("Current collector perpendicular area [m2]")

5.7.3.2 A_{cc} ep_bolfi.models.standard_parameters. A_{cc} = A

5.7.3.3 a_n ep_bolfi.models.standard_parameters. a_n = $a_{n_dim} * R_n$

5.7.3.4 a_{n_dim} ep_bolfi.models.standard_parameters. a_{n_dim} = Parameter("Negative electrode surface area to volume ratio [m-1]")

5.7.3.5 a_p ep_bolfi.models.standard_parameters. a_p = $a_{p_dim} * R_p$

5.7.3.6 a_{p_dim} ep_bolfi.models.standard_parameters. a_{p_dim} = Parameter("Positive electrode surface area to volume ratio [m-1]")

5.7.3.7 C ep_bolfi.models.standard_parameters. C = Parameter("Typical current [A]")

5.7.3.8 capacity `ep_bolfi.models.standard_parameters.capacity = Parameter("Nominal cell capacity [A.h]")`

5.7.3.9 current_with_time `ep_bolfi.models.standard_parameters.current_with_time = I_extern`

5.7.3.10 C_{rn} `ep_bolfi.models.standard_parameters.Crn = τrn / τd`

5.7.3.11 C_{rp} `ep_bolfi.models.standard_parameters.Crp = τrp / τd`

5.7.3.12 C_e `int ep_bolfi.models.standard_parameters.Ce = τe / τd`

5.7.3.13 $c_e_dim_init$ `ep_bolfi.models.standard_parameters.ce_dim_init = Parameter("Initial concentration in electrolyte [mol.↔m-3]")`

5.7.3.14 c_e_init `ep_bolfi.models.standard_parameters.ce_init = ce_dim_init / ce_typ`

5.7.3.15 c_e_typ `ep_bolfi.models.standard_parameters.ce_typ = pybamm.Parameter("Typical electrolyte concentration [mol.m-3]")`

5.7.3.16 c_n `ep_bolfi.models.standard_parameters.cn = pybamm.Parameter("Maximum concentration in negative electrode [mol.↔m-3]")`

5.7.3.17 C_n `int ep_bolfi.models.standard_parameters.Cn = τn / τd`

5.7.3.18 c_p `ep_bolfi.models.standard_parameters.cp = pybamm.Parameter("Maximum concentration in positive electrode [mol.m-3]")`

5.7.3.19 C_p `int ep_bolfi.models.standard_parameters.C_p = τ_p / τ^d`

5.7.3.20 `dimensional_current_density_with_time` `ep_bolfi.models.standard_parameters.dimensional_current_density_with_time = $I_{\text{extern_dim}} / A$`

5.7.3.21 `dimensional_current_with_time` `ep_bolfi.models.standard_parameters.dimensional_current_with_time = $I_{\text{extern_dim}}$`

5.7.3.22 D_e _typ `def ep_bolfi.models.standard_parameters.D_e_typ = $D_{e_dim}(c_{e_typ}, T_{\text{ref}})$`

5.7.3.23 D_n _typ `def ep_bolfi.models.standard_parameters.D_n_typ = $D_{n_dim}(\text{Scalar}(0.5), T_{\text{ref}})$`

5.7.3.24 D_p _typ `def ep_bolfi.models.standard_parameters.D_p_typ = $D_{p_dim}(\text{Scalar}(0.5), T_{\text{ref}})$`

5.7.3.25 F `ep_bolfi.models.standard_parameters.F = $\text{Scalar}(\text{constants.physical_constants}["\text{Faraday constant}"][0])$`

5.7.3.26 I_{extern} `ep_bolfi.models.standard_parameters.I_extern = $I_{\text{extern_dim}} / C$`

5.7.3.27 $I_{\text{extern_dim}}$ `ep_bolfi.models.standard_parameters.I_extern_dim`

Initial value:

```
1 = pybamm.FunctionParameter(
2     "Current function [A]",
3     {"Time [s]": pybamm.t * timescale}
4 )
```

5.7.3.28 I_{typ} `ep_bolfi.models.standard_parameters.I_typ = C`

5.7.3.29 $i_{sn_0_ref}$ `def ep_bolfi.models.standard_parameters.isn_0_ref = isn_0_dim(ce_typ, 0.5 * cn, cn, T_ref)`

5.7.3.30 $i_{sep_0_ref}$ `def ep_bolfi.models.standard_parameters.isep_0_ref = isep_0_dim(ce_typ, 0.5 * cp, cp, T_ref)`

5.7.3.31 k_B `ep_bolfi.models.standard_parameters.k_B = constants.physical_constants["Boltzmann constant"][0]`

5.7.3.32 L_dim `ep_bolfi.models.standard_parameters.L_dim = Ln_dim + Ls_dim + Lp_dim`

5.7.3.33 L_x `ep_bolfi.models.standard_parameters.L_x = L_dim`

5.7.3.34 L_y `ep_bolfi.models.standard_parameters.L_y = Parameter("Electrode width [m]")`

5.7.3.35 L_z `ep_bolfi.models.standard_parameters.L_z = Parameter("Electrode height [m]")`

5.7.3.36 L_e `ep_bolfi.models.standard_parameters.Le`

Initial value:

```
1 = pybamm.Concatenation(
2   pybamm.PrimaryBroadcast(Ln, "negative electrode"),
3   pybamm.PrimaryBroadcast(Ls, "separator"),
4   pybamm.PrimaryBroadcast(Lp, "positive electrode")
5 )
```

5.7.3.37 L_n `ep_bolfi.models.standard_parameters.Ln = Ln_dim / L_dim`

5.7.3.38 L_n_dim `ep_bolfi.models.standard_parameters.Ln_dim = Parameter("Negative electrode thickness [m]")`

5.7.3.39 **L_p** ep_bolfi.models.standard_parameters. L_p = L_{p_dim} / L_dim

5.7.3.40 **L_{p_dim}** ep_bolfi.models.standard_parameters. L_{p_dim} = Parameter("Positive electrode thickness [m]")

5.7.3.41 **L_s** ep_bolfi.models.standard_parameters. L_s = L_{s_dim} / L_dim

5.7.3.42 **L_{s_dim}** ep_bolfi.models.standard_parameters. L_{s_dim} = Parameter("Separator thickness [m]")

5.7.3.43 **n_cells** ep_bolfi.models.standard_parameters. n_cells = Parameter("Number of cells connected in series to make a battery")

5.7.3.44 **$n_electrodes_parallel$** ep_bolfi.models.standard_parameters. $n_electrodes_parallel$

Initial value:

1 = Parameter("Number of electrodes connected in parallel"
2 " to make a cell")

5.7.3.45 **OCV_{n_ref}** def ep_bolfi.models.standard_parameters. OCV_{n_ref} = $OCV_{n_dim}(SOC_{n_init}(0), T_ref)$

5.7.3.46 **OCV_{p_ref}** def ep_bolfi.models.standard_parameters. OCV_{p_ref} = $OCV_{p_dim}(SOC_{p_init}(1), T_ref)$

5.7.3.47 **Q** tuple ep_bolfi.models.standard_parameters. Q = $(1 - \epsilon_{p_scalar}) * L_{p_dim} * c_p * z_p * F * A$

5.7.3.48 **q_e** ep_bolfi.models.standard_parameters. q_e = constants.physical_constants["electron volt"][0]

5.7.3.49 **R** ep_bolfi.models.standard_parameters. R = Scalar(constants.R)

5.7.3.50 R_n `ep_bolfi.models.standard_parameters.R_n = Parameter("Negative particle radius [m]")`

5.7.3.51 R_p `ep_bolfi.models.standard_parameters.R_p = Parameter("Positive particle radius [m]")`

5.7.3.52 T_{init} `ep_bolfi.models.standard_parameters.T_init = Parameter("Initial temperature [K]")`

5.7.3.53 T_{ref} `ep_bolfi.models.standard_parameters.T_ref = Parameter("Reference temperature [K]")`

5.7.3.54 $thermal_voltage$ `ep_bolfi.models.standard_parameters.thermal_voltage = $R * T_{ref} / F$`

5.7.3.55 $timescale$ `ep_bolfi.models.standard_parameters.timescale = τ^d`

5.7.3.56 U_u `ep_bolfi.models.standard_parameters.U_u = pybamm.Parameter("Upper voltage cut-off [V]")`

5.7.3.57 U_l `ep_bolfi.models.standard_parameters.U_l = pybamm.Parameter("Lower voltage cut-off [V]")`

5.7.3.58 V `ep_bolfi.models.standard_parameters.V = Parameter("Cell volume [m3]")`

5.7.3.59 $voltage_high_cut$ `ep_bolfi.models.standard_parameters.voltage_high_cut = U_u`

5.7.3.60 $voltage_low_cut$ `ep_bolfi.models.standard_parameters.voltage_low_cut = U_l`

5.7.3.61 z_n `ep_bolfi.models.standard_parameters.z_n = Parameter("Negative electrode electrons in reaction")`

5.7.3.62 z_p ep_bolfi.models.standard_parameters. z_p = Parameter("Positive electrode electrons in reaction")

5.7.3.63 ΔT ep_bolfi.models.standard_parameters. ΔT = Scalar(1)

5.7.3.64 α_{mn} ep_bolfi.models.standard_parameters. α_{mn} = Parameter("Negative electrode anodic charge-transfer coefficient")

5.7.3.65 α_{np} ep_bolfi.models.standard_parameters. α_{np} = Parameter("Positive electrode anodic charge-transfer coefficient")

5.7.3.66 α_{pn} ep_bolfi.models.standard_parameters. α_{pn} = Parameter("Negative electrode cathodic charge-transfer coefficient")

5.7.3.67 α_{pp} ep_bolfi.models.standard_parameters. α_{pp} = Parameter("Positive electrode cathodic charge-transfer coefficient")

5.7.3.68 β_e ep_bolfi.models.standard_parameters. β_e = pybamm.Concatenation(β_a , β_{es} , β_{ep})

5.7.3.69 β_a ep_bolfi.models.standard_parameters. β_a = pybamm.PrimaryBroadcast(β_a_scalar , "negative electrode")

5.7.3.70 β_a_scalar ep_bolfi.models.standard_parameters. β_a_scalar

Initial value:

```
1 = Parameter(
2   "Negative electrode Bruggeman coefficient (electrolyte)"
3 )
```

5.7.3.71 β_{ep} ep_bolfi.models.standard_parameters. β_{ep} = pybamm.PrimaryBroadcast(β_{ep_scalar} , "positive electrode")

5.7.3.72 β_{ep_scalar} `ep_bolfi.models.standard_parameters. β_{ep_scalar}`

Initial value:

```
1 = Parameter(
2   "Positive electrode Bruggeman coefficient (electrolyte)"
3 )
```

5.7.3.73 β_{es} `ep_bolfi.models.standard_parameters. β_{es} = pybamm.PrimaryBroadcast(β_{es_scalar} , "separator")`

5.7.3.74 β_{es_scalar} `ep_bolfi.models.standard_parameters. β_{es_scalar} = Parameter("Separator Bruggeman coefficient (electrolyte)")`

5.7.3.75 β_{sn_scalar} `ep_bolfi.models.standard_parameters. β_{sn_scalar} = Parameter("Negative electrode Bruggeman coefficient (electrode)")`

5.7.3.76 β_{sp_scalar} `ep_bolfi.models.standard_parameters. β_{sp_scalar} = Parameter("Positive electrode Bruggeman coefficient (electrode)")`

5.7.3.77 β_{ss_scalar} `ep_bolfi.models.standard_parameters. β_{ss_scalar} = Parameter("Separator Bruggeman coefficient (electrode)")`

5.7.3.78 γ_e `ep_bolfi.models.standard_parameters. γ_e = c_{e_typ} / c_p`

5.7.3.79 γ_n `ep_bolfi.models.standard_parameters. γ_n = c_n / c_p`

5.7.3.80 γ_p `ep_bolfi.models.standard_parameters. γ_p = c_p / c_p`

5.7.3.81 ϵ `ep_bolfi.models.standard_parameters. ϵ = pybamm.Concatenation(ϵ_n , ϵ_s , ϵ_p)`

5.7.3.82 ϵ^β ep_bolfi.models.standard_parameters. $\epsilon^\beta = \epsilon ** \beta_e$

5.7.3.83 ϵ_n ep_bolfi.models.standard_parameters. $\epsilon_n = \text{pybamm.PrimaryBroadcast}(\epsilon_n_scalar, \text{"negative electrode"})$

5.7.3.84 ϵ_n_scalar ep_bolfi.models.standard_parameters. $\epsilon_n_scalar = \text{Parameter}(\text{"Negative electrode porosity"})$

5.7.3.85 ϵ_p ep_bolfi.models.standard_parameters. $\epsilon_p = \text{pybamm.PrimaryBroadcast}(\epsilon_p_scalar, \text{"positive electrode"})$

5.7.3.86 ϵ_p_scalar ep_bolfi.models.standard_parameters. $\epsilon_p_scalar = \text{Parameter}(\text{"Positive electrode porosity"})$

5.7.3.87 ϵ_s ep_bolfi.models.standard_parameters. $\epsilon_s = \text{pybamm.PrimaryBroadcast}(\epsilon_s_scalar, \text{"separator"})$

5.7.3.88 ϵ_s_scalar ep_bolfi.models.standard_parameters. $\epsilon_s_scalar = \text{Parameter}(\text{"Separator porosity"})$

5.7.3.89 κ_e_hat tuple ep_bolfi.models.standard_parameters. $\kappa_e_hat = (R * T_ref / F) / (C / A * L_dim / \kappa_e_typ)$

5.7.3.90 κ_e_typ def ep_bolfi.models.standard_parameters. $\kappa_e_typ = \kappa_e_dim(c_e_typ, T_ref)$

5.7.3.91 σ_n tuple ep_bolfi.models.standard_parameters. $\sigma_n = (\text{thermal_voltage} / (C / A * L_dim)) * \sigma_n_dim$

5.7.3.92 σ_n_dim ep_bolfi.models.standard_parameters. $\sigma_n_dim = \text{pybamm.Parameter}(\text{"Negative electrode conductivity [S.m-1]"})$

5.7.3.93 σ_p tuple ep_bolfi.models.standard_parameters. $\sigma_p = (\text{thermal_voltage} / (C / A * L_dim)) * \sigma_p_dim$

5.7.3.94 σ_{p_dim} `ep_bolfi.models.standard_parameters. σ_{p_dim} = pybamm.Parameter("Positive electrode conductivity [S.m-1]")`

5.7.3.95 τ^d `ep_bolfi.models.standard_parameters. τ^d = $F * c_p * L_{dim} / (C / A)$`

5.7.3.96 τ_{rn} `ep_bolfi.models.standard_parameters. τ_{rn} = $F * c_n / (i_{se0_ref} * a_{n_dim})$`

5.7.3.97 τ_{rp} `ep_bolfi.models.standard_parameters. τ_{rp} = $F * c_p / (i_{sep0_ref} * a_{p_dim})$`

5.7.3.98 τ_e `int ep_bolfi.models.standard_parameters. τ_e = $L_{dim} ** 2 / D_{e_typ}$`

5.7.3.99 τ_n `int ep_bolfi.models.standard_parameters. τ_n = $R_n ** 2 / D_{n_typ}$`

5.7.3.100 τ_p `int ep_bolfi.models.standard_parameters. τ_p = $R_p ** 2 / D_{p_typ}$`

5.8 `ep_bolfi.optimization` Namespace Reference

Namespaces

- [EP_BOLFI](#)

5.9 `ep_bolfi.optimization.EP_BOLFI` Namespace Reference

Classes

- class [NDArrayEncoder](#)
- class [Preprocessed_Simulator](#)
Normalizes sampling to a standard normal distribution.
- class [Optimizer_State](#)
Handles the heuristics for the EP-BOLFI operation modes.
- class [EP_BOLFI](#)
Expectation Propagation and Bayesian Optimization.

Functions

- def [combine_parameters_to_try](#) (parameters, parameters_to_try_dict)
Give every combination as full parameter sets.
- def [fix_parameters](#) (parameters_to_be_fixed)
Returns a function which sets some parameters in advance.

5.9.1 Function Documentation

5.9.1.1 combine_parameters_to_try() def ep_bolfi.optimization.EP_BOLFI.combine_parameters_to_try (
parameters,
parameters_to_try_dict)

Give every combination as full parameter sets.

Parameters

<i>parameters</i>	The base full parameter set as a dictionary.
<i>parameters_to_try_dict</i>	The keys of this dictionary correspond to the "parameters" keys where different values are to be inserted. These are given by the tuples which are the values of this dictionary.

Returns

A 2-tuple where the first item is the list of all parameter set combinations and the second the list of the combinations only.

5.9.1.2 fix_parameters() def ep_bolfi.optimization.EP_BOLFI.fix_parameters (
parameters_to_be_fixed)

Returns a function which sets some parameters in advance.

Parameters

<i>parameters_to_be_fixed</i>	These parameters will at least be a part of the dictionary that the returned function returns.
-------------------------------	--

Returns

The function which adds additional parameters to a dictionary or replaces existing parameters with the new ones.

5.10 ep_bolfi.utility Namespace Reference

Namespaces

- [dataset_formatting](#)

- *Defines datatypes for further processing.*
- [fitting_functions](#)
Various helper and fitting functions for processing measurement curves.
- [preprocessing](#)
Contains frequently used workflows in dataset preprocessing.
- [visualization](#)
Various helper and plotting functions for common data visualizations.

5.11 ep_bolfi.utility.dataset_formatting Namespace Reference

Defines datatypes for further processing.

Classes

- class [Measurement](#)
Defines common methods for measurement objects.
- class [Cycling_Information](#)
Contains basic cycling informations.
- class [Static_Information](#)
Contains additional informations, e.g.
- class [Impedance_Measurement](#)
Contains basic impedance data.

Functions

- def [read_csv_from_measurement_system](#) (path, encoding, number_of_comment_lines, headers, delimiter='\\t', decimal='.', datatype="cycling", segment_column=-1, segments_to_process=None, current_sign_correction={}, correction_column=-1, flip_voltage_sign=False, flip_imaginary_impedance_sign=False, max_number_of_lines=-1)
Read the measurements as returned by common instruments.
- def [print_hdf5_structure](#) (h5py_object, depth=1, table_limit=4, verbose_limit=64)
brief Simple HDF5 structure viewer.
- def [convert_none_notation_to_slicing](#) (h5py_object, index)
Access a slice of an HDF5 object by transferable notation.
- def [get_hdf5_dataset_by_path](#) (h5py_object, path)
Follow the structure of a HDF object to get a certain part.
- def [read_hdf5_table](#) (path, data_location, headers, datatype="cycling", segment_location=None, segments_to_process=None, current_sign_correction={}, correction_location=None, flip_voltage_sign=False, flip_imaginary_impedance_sign=False)
Read the measurements as stored in a HDF5 file.

5.11.1 Detailed Description

Defines datatypes for further processing.

5.11.2 Function Documentation

5.11.2.1 [convert_none_notation_to_slicing\(\)](#) def ep_bolfi.utility.dataset_formatting.convert_none_notation_to_slicing (h5py_object, index)

Access a slice of an HDF5 object by transferable notation.

Parameters

<i>h5py_object</i>	The HDF5 object wrapped by H5Py, for example <code>h5py.File(filename, 'r')</code> .
<i>index</i>	A 2-tuple or a 2-list. (None, x) denotes slicing <code>[:, x]</code> and (x, None) denotes slicing <code>[x, :]</code> .

5.11.2.2 get_hdf5_dataset_by_path() `def ep_bolfi.utility.dataset_formatting.get_hdf5_dataset_by_path (`
h5py_object,
path
`)`

Follow the structure of a HDF object to get a certain part.

Parameters

<i>h5py_object</i>	The HDF5 object wrapped by H5Py, for example <code>h5py.File(filename, 'r')</code> .
<i>path</i>	A list. Each entry goes one level deeper into the HDF structure. Each entry can either be the index to go into next itself, or a 2-tuple or a 2-list. In the latter case, (None, x) denotes slicing <code>[:, x]</code> and (x, None) denotes slicing <code>[x, :]</code> .

Returns

Returns the HDF5 object found at the end of "path".

5.11.2.3 print_hdf5_structure() `def ep_bolfi.utility.dataset_formatting.print_hdf5_structure (`
h5py_object,
depth = 1,
table_limit = 4,
verbose_limit = 64)
`)`

brief Simple HDF5 structure viewer.

Parameters

<i>h5py_object</i>	The HDF5 object wrapped by H5Py, for example <code>h5py.File(filename, 'r')</code> .
<i>depth</i>	For pretty printing the recursive depth. Do not change.
<i>table_limit</i>	<code>h5py.Dataset</code> object tables will be truncated prior to printing up to this number in the higher dimension.
<i>verbose_limit</i>	<code>h5py.Dataset</code> objects will be truncated to at most this number in any dimension.

5.11.2.4 read_csv_from_measurement_system() `def ep_bolfi.utility.dataset_formatting.read_csv_from_measurement←↵`
`_system (`
path,
encoding,
`)`


```

number_of_comment_lines,
headers,
delimiter = '\t',
decimal = '.',
datatype = "cycling",
segment_column = -1,
segments_to_process = None,
current_sign_correction = {},
correction_column = -1,
flip_voltage_sign = False,
flip_imaginary_impedance_sign = False,
max_number_of_lines = -1 )

```

Read the measurements as returned by common instruments.

Example: cycling measurements from Basytec devices. Their format resembles a csv file with one title and one header comment line. So the first line will be ignored and the second used for headers.

Parameters

<i>path</i>	The full or relative path to the measurement file.
<i>encoding</i>	The encoding of that file, e.g. "iso-8859-1".
<i>number_of_comment_lines</i>	The number of lines that have to be skipped over in order to arrive at the first dataset line.
<i>headers</i>	A dictionary. Its keys are the indices of the columns which are to be read in. The corresponding values are there to tell this function which kind of data is in which column. The following format has to be used: "<name> [<unit>]" where "name" is "U" (voltage), "I" (current) or "t" (time) and "unit" is "V", "A", "h", "m" or "s" with the optional prefixes "k", "m", "μ" or "n". This converts the data to prefix-less SI units. Additional columns may be read in with keys not in this format. The columns for segments and sign correction are only given by #segment_column and #correction_column.
<i>delimiter</i>	The delimiter string between datapoints. The default is "\t".
<i>decimal</i>	The string used for the decimal point. Default: ".".
<i>datatype</i>	Default is "cycling", where cycling information is assumed in the file. "static" will trigger the additional extraction of exponential decays that are relevant to e.g. GITT. "impedance" will treat the file as an impedance measurement with frequencies and impedances instead of time and voltage.
<i>segment_column</i>	The index of the column that stores the index of the current segment. If it changes from one data point to the next, that is used as the dividing line between two segments. Default is -1, which returns the dataset in one segment.
<i>segments_to_process</i>	A list of indices which give the segments that shall be processed. Default is None, i.e., the whole file gets processed.
<i>current_sign_correction</i>	A dictionary. Its keys are the strings used in the file to indicate a state. The column from which this state is retrieved is given by #correction_column. The dictionaries' values are used to correct/normalize the current value in the file. For example, if discharge currents have the same positive sign as charge currents in the file, use -1 to correct that, or if the values are to be scaled by weight, use the scaling factor. The default is the empty dictionary.
<i>correction_column</i>	See #current_sign_correction. Default: -1.
<i>max_number_of_lines</i>	The maximum number of dataset lines that are to be read in. Default: -1 (no limit).

Parameters

<i>flip_voltage_sign</i>	Defaults to False, where measured voltage remains unaltered. Change to True if the voltage shall be multiplied by -1. Also applies to impedances; real and imaginary parts.
<i>flip_imaginary_impedance_sign</i>	Defaults to False, where measured impedance remains unaltered. Change to True if the imaginary part of the impedance shall be multiplied by -1. Cancels out with 'flip-voltage-sign'.

Returns

The measurement, packaged in a [Measurement](#) subclass. It depends on "datatype" which one it is:

- "cycling": [Cycling_Information](#)
- "static": [Static_Information](#)
- "impedance": [Impedance_Measurement](#)

5.11.2.5 read_hdf5_table() `def ep_bolfi.utility.dataset_formatting.read_hdf5_table (`
path,
data_location,
headers,
datatype = "cycling",
segment_location = None,
segments_to_process = None,
current_sign_correction = {},
correction_location = None,
flip_voltage_sign = False,
flip_imaginary_impedance_sign = False)

Read the measurements as stored in a HDF5 file.

Parameters

<i>path</i>	The full or relative path to the measurement file.
<i>data_location</i>	A list. Gives the location in the HDF5 file where the data table is stored. Set to None if everything is stored at the top level. Each entry goes one level deeper into the HDF structure. Each entry can either be the index to go into next itself, or a 2-tuple or a 2-list. In the latter case, (None, x) denotes slicing [:, x] and (x, None) denotes slicing [x, :].
<i>headers</i>	A dictionary. Its keys are 2-tuples, slicing the data which are to be read in. Use the format "(x, None)" or "(None, x)" to slice the dimension with "None". The corresponding values are there to tell this function which kind of data is in which column. If necessary, the keys may also be tuples like 'data_location'. The following format has to be used: "<name> [<unit>]" where "name" is "U" (voltage), "I" (current) or "t" (time) and "unit" is "V", "A", "h", "m" or "s" with the optional prefixes "k", "m", "μ" or "n". This converts the data to prefix-less SI units. Additional columns may be read in with keys not in this format. The columns for segments and sign correction are only given by "segment_column" and "correction_column".

Parameters

<i>datatype</i>	Default is "cycling", where cycling information is assumed in the file. "static" will trigger the additional extraction of exponential decays that are relevant to e.g. GITT. "impedance" will treat the file as an impedance measurement with frequencies and impedances instead of time and voltage. @segment_location A list, with the same format as "data_location". It points to the part of the data that stores the index of the current segment. If it changes from one data point to the next, that is used as the dividing line between two segments. Default is None, which returns the dataset in one segment.
<i>segments_to_process</i>	A list of indices which give the segments that shall be processed. Default is None, i.e., the whole file gets processed.
<i>current_sign_correction</i>	A dictionary. Its keys are the strings used in the file to indicate a state. The column from which this state is retrieved is given by "correction_column". The dictionaries' values are used to correct/normalize the current value in the file. For example, if discharge currents have the same positive sign as charge currents in the file, use -1 to correct that, or if the values are to be scaled by weight, use the scaling factor. The default is the empty dictionary.
<i>correction_location</i>	A list, with the same format as "data_location". For its use, see "current_sign_correction". Default: None.
<i>flip_voltage_sign</i>	Defaults to False, where measured voltage remains unaltered. Change to True if the voltage shall be multiplied by -1. Also applies to impedances; real and imaginary parts.
<i>flip_imaginary_impedance_sign</i>	Defaults to False, where measured impedance remains unaltered. Change to True if the imaginary part of the impedance shall be multiplied by -1. Cancels out with 'flip-voltage-sign'.

Returns

The measurement, packaged in a [Measurement](#) subclass. It depends on "datatype" which one it is:

- "cycling": [Cycling_Information](#)
- "static": [Static_Information](#)
- "impedance": [Impedance_Measurement](#)

5.12 ep_bolfi.utility.fitting_functions Namespace Reference

Various helper and fitting functions for processing measurement curves.

Classes

- class [NDArrayEncoder](#)
- class [OCV_fit_result](#)

Contains OCV fit parameters and related information.

Functions

- def [find_occurrences](#) (sequence, value)
Gives indices in sequence where it is closest to value.
- def [smooth_fit](#) (x, y, order=3, splits=None, w=None, s=None, display=False, derivatives=0)
Calculates a smoothed spline with derivatives.
- def [fit_exponential_decay](#) (timepoints, voltages, recursive_depth=1, threshold=0.95)
Extracts a set amount of exponential decay curves.
- def [fit_sqrt](#) (timepoints, voltages, threshold=0.95)
Extracts a square root at the beginning of the data.
- def [fit_drt](#) (frequencies, impedances, lambda_value=-2.0)
- def [laplace_transform](#) (x, y, s)
Performs a basic laplace transformation.
- def [a_fit](#) (yUminus1)
Calculates the conversion from "A parametric OCV model".
- def [OCV_fit_function](#) (E_OCV, *args, z=1.0, T=298.15, individual=False, fit_SOC_range=False, rescale=False)
The OCV model from "A parametric OCV model".
- def [d_dE_OCV_fit_function](#) (E_OCV, *args, z=1.0, T=298.15, individual=False, fit_SOC_range=False, rescale=False)
The derivative of [fitting_functions.OCV_fit_function](#).
- def [d2_dE2_OCV_fit_function](#) (E_OCV, *args, z=1.0, T=298.15, individual=False, fit_SOC_range=False, rescale=False)
The 2ⁿ derivative of [fitting_functions.OCV_fit_function](#).
- def [inverse_OCV_fit_function](#) (SOC, *args, z=1.0, T=298.15, inverted=True)
The inverse of [fitting_functions.OCV_fit_function](#).
- def [inverse_d_dSOC_OCV_fit_function](#) (SOC, *args, z=1.0, T=298.15, inverted=True)
The derivative of the inverse of [fitting_functions.OCV_fit_function](#).
- def [inverse_d2_dSOC2_OCV_fit_function](#) (SOC, *args, z=1.0, T=298.15, inverted=True)
The 2nd derivative of the inverse of [fitting_functions.OCV_fit_function](#).
- def [fit_OCV](#) (SOC, OCV, N=4, SOC_range_bounds=(0.2, 0.8), SOC_range_limits=(0.0, 1.0), z=1.0, T=298.15, inverted=True, fit_SOC_range=True, distance_order=2, weights=None, initial_parameters=None, minimize_options=None)
Fits data to [fitting_functions.OCV_fit_function](#).
- def [verbose_spline_parameterization](#) (coeffs, knots, order, format='python', function_name="OCV", function_args="SOC", derivatives=0, spline_transformation="", verbose=False)
Gives the monomic representation of a B-spline.

5.12.1 Detailed Description

Various helper and fitting functions for processing measurement curves.

5.12.2 Function Documentation

5.12.2.1 a_fit() def ep_bolfi.utility.fitting_functions.a_fit (yUminus1)

Calculates the conversion from "A parametric OCV model".

Parameters

γU_{minus1}	$\gamma * U_i / e$ from "A parametric OCV model".
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Returns

The approximation factor a_i from "A parametric OCV model".

5.12.2.2 d2_dE2_OCV_fit_function() `def ep_bolfi.utility.fitting_functions.d2_dE2_OCV_fit_function (`
`E_OCV,`
`* args,`
`z = 1.0,`
`T = 298.15,`
`individual = False,`
`fit_SOC_range = False,`
`rescale = False)`

The 2nd derivative of [fitting_functions.OCV_fit_function](#).

Parameters

<i>E_OCV</i>	The voltages for which the 2 nd derivative shall be evaluated.
<i>args</i>	A list which length is dividable by three. These are the parameters E_0 , a and Δx from "A parametric OCV model" in the order (E_{0_0} , a_{0_0} , Δx_{0_0} , E_{0_1} , a_{0_1} , Δx_{0_1} ...). The last Δx_j may be omitted to force $\sum \Delta x_j = 1$.
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>individual</i>	If True, the model function summands are not summed up.
<i>fit_SOC_range</i>	If True, this function takes two additional arguments at the start of "args" that may be used to adjust the data SOC range.
<i>rescale</i>	If True, the expected "args" now contain the slopes of the summands at their respective origins instead of a . Formula: $a / \text{slope} = -4 * (k_B / e) * T / (\Delta x * z)$.

Returns

The evaluation of $\partial^2 \text{OCV_fit_function}(\text{OCV}) / \partial^2 \text{OCV}$.

5.12.2.3 d_dE_OCV_fit_function() `def ep_bolfi.utility.fitting_functions.d_dE_OCV_fit_function (`
`E_OCV,`
`* args,`
`z = 1.0,`
`T = 298.15,`
`individual = False,`
`fit_SOC_range = False,`
`rescale = False)`

The derivative of [fitting_functions.OCV_fit_function](#).

Parameters

<i>E_OCV</i>	The voltages for which the derivative shall be evaluated.
<i>args</i>	A list which length is dividable by three. These are the parameters E_0 , a and Δx from "A parametric OCV model" in the order (E_{0_0} , a_{0_0} , Δx_{0_0} , E_{0_1} , a_{0_1} , Δx_{0_1} ...). The last Δx_j may be omitted to force $\sum \Delta x_j = 1$.
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>individual</i>	If True, the model function summands are not summed up.
<i>fit_SOC_range</i>	If True, this function takes two additional arguments at the start of "args" that may be used to adjust the data SOC range.
<i>rescale</i>	If True, the expected "args" now contain the slopes of the summands at their respective origins instead of a . Formula: $a / \text{slope} = -4 * (k_B / e) * T / (\Delta x * z)$.

Returns

The evaluation of $\partial \text{OCV_fit_function}(\text{OCV}) / \partial \text{OCV}$.

5.12.2.4 find_occurrences() def ep_bolfi.utility.fitting_functions.find_occurrences (
sequence,
value)

Gives indices in sequence where it is closest to value.

Parameters

<i>sequence</i>	A list that represents a differentiable function.
<i>value</i>	The value that is searched for in "sequence". Also, crossings of consecutive values in "sequence" with "value" are searched for.

Returns

A list of indices in "sequence" in ascending order where "value" or a close match for "value" was found.

5.12.2.5 fit_drt() def ep_bolfi.utility.fitting_functions.fit_drt (
frequencies,
impedances,
lambda_value = -2.0)

Distribution of Relaxation Times.

@param frequencies

Array of the measured frequencies.

@param impedances

Array of the measured complex impedances.

@param lambda_value

Takes the place of the R^2 value as a tuning parameter.

Consult the pyimpspec documentation for the precise usage.

Default is -2, which "uses the L-curve approach to estimate lambda".
 -1 uses a different heuristic, and values > 0 set lambda directly.

@return

A 3-tuple of characteristic DRT time constants, their corresponding resistances, and the whole TRNNLSResult object returned by pyimpspec.

Note that the .pseudo_chisqr attribute is not useful for the same reasons that went into the R^2 calculations in the other fit functions.

5.12.2.6 fit_exponential_decay() `def ep_bolfi.utility.fitting_functions.fit_exponential_decay (`
 timepoints,
 voltages,
 recursive_depth = 1,
 threshold = 0.95)

Extracts a set amount of exponential decay curves.

Parameters

<i>timepoints</i>	The timepoints of the measurements.
<i>voltages</i>	The corresponding voltages.
<i>recursive_depth</i>	The default (1) fits one exponential curve to the data. For higher values that fit is repeated with the data minus the preceding fit(s) for this amount of times minus one.
<i>threshold</i>	The lower threshold value for the R^2 coefficient of determination. If "threshold" is smaller than 1, the subset of the exponential decay data is searched that just fulfills it. Defaults to 0.95. Values above 1 are set to 1.

Returns

A list of length "recursive_depth" where each element is a 3-tuple with the timepoints, the fitted voltage evaluations and a 3-tuple of the parameters of the following decay function: $t, (U_0, \Delta U, \tau_r^{-1}) \mapsto U_0 + \Delta U * \exp(-\tau_r^{-1} * (t - \text{timepoints}[0]))$.

5.12.2.7 fit_OCV() `def ep_bolfi.utility.fitting_functions.fit_OCV (`
 SOC,
 OCV,
 N = 4,
 SOC_range_bounds = (0.2, 0.8),
 SOC_range_limits = (0.0, 1.0),
 z = 1.0,
 T = 298.15,
 inverted = True,
 fit_SOC_range = True,
 distance_order = 2,
 weights = None,
 initial_parameters = None,
 minimize_options = None)

Fits data to [fitting_functions.OCV_fit_function](#).

In addition to the fit itself, a model-based correction to the provided SOC-OCV-data is made. If "SOC" lives in a (0,1)-range, the correction is given as its transformation to the (SOC_start, SOC_end)-range given as the first two returned numbers. If "other_electrode" is not None, the stoichiometric offset and the scaling that gives the (adjusted) SOC of this electrode given the SOC of the "other_electrode" are put before SOC_start and SOC_end.

Parameters

<i>SOC</i>	The SOC's at which measurements were made.
<i>OCV</i>	The corresponding open-circuit voltages.
<i>N</i>	The number of phases of the OCV model.
<i>SOC_range_bounds</i>	Optional hard upper and lower bounds for the SOC correction from the left and the right side, respectively, as a 2-tuple. Use it as a limiting guess for the actual SOC range represented in the measurement. Has to be inside (0.0, 1.0). Set to (0.0, 1.0) to effectively disable SOC range estimation.
<i>SOC_range_limits</i>	Optional hard lower and upper bounds for the SOC correction from the left and the right side, respectively, as a 2-tuple. Use it if you know that your OCV data is incomplete and by how much. Has to be inside (0.0, 1.0). Set to (0.0, 1.0) to allow the SOC range estimation to assign datapoints to the asymptotes.
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>inverted</i>	If True (default), the widely adopted SOC convention is assumed. If False, the formulation of "A parametric OCV model" is used.
<i>fit_SOC_range</i>	If True (default), a model-based correction to the provided SOC-OCV-data is made.
<i>distance_order</i>	The order of the norm of the vector of the distances between OCV data and OCV model. Default is 2, i.e., the Euclidean norm. 1 sets it to absolute distance, and float('inf') sets it to maximum distance. Note that 1 will lead to worse performance.
<i>weights</i>	Optional weights to apply to the vector of the distances between OCV data and OCV model. Defaults to equal weights.
<i>initial_parameters</i>	Optional initial guess for the model parameters. If left as-is, this will be automatically gleaned from the data. Use only if you have another fit to data of the same electrode material.
<i>minimize_options</i>	Dictionary that gets passed to <code>scipy.optimize.minimize</code> with the method 'trust-constr'. See <code>scipy.optimize.show_options</code> with the arguments 'minimize' and 'trust-constr' for details.

Returns

The fitted parameters of [fitting_functions.OCV_fit_function](#) plus the fitted SOC range prepended.

5.12.2.8 `fit_sqrt()` `def ep_bolfi.utility.fitting_functions.fit_sqrt (`
 timepoints,
 voltages,
 threshold = 0.95)

Extracts a square root at the beginning of the data.

Parameters

<i>timepoints</i>	The timepoints of the measurements.
<i>voltages</i>	The corresponding voltages.
<i>threshold</i>	The lower threshold value for the R^2 coefficient of determination. If "threshold" is smaller than 1, the subset of the experimental data is searched that just fulfills it. Defaults to 0.95. Values above 1 are set to 1.

Returns

A 3-tuple with the timepoints, the fitted voltage evaluations and a 2-tuple of the parameters of the following sqrt function: $t, (U_0, dU_d\sqrt{t}) \mapsto U_0 + dU_d\sqrt{t} * \sqrt{(t - \text{timepoints}[0])}$.

5.12.2.9 inverse_d2_dSOC2_OCV_fit_function() `def ep_bolfi.utility.fitting_functions.inverse_d2_dSOC2_OCV_fit_function (`
`SOC,`
`* args,`
`z = 1.0,`
`T = 298.15,`
`inverted = True)`

The 2nd derivative of the inverse of .OCV_fit_function.

Basically OCV'(SOC). Requires that the Δx entries are sorted by x. This corresponds to the parameters being sorted by decreasing E_0 .

Parameters

<i>E_OCV</i>	The SOC's for which the voltages shall be evaluated.
<i>args</i>	A list which length is dividable by three. These are the parameters E_0 , a and Δx from "A parametric OCV model" in the order (E_{0_0} , a_0 , Δx_0 , E_{0_1} , a_1 , $\Delta x_1...$).
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>inverted</i>	The default (False) uses the formulation from "A parametric OCV model". If True, the SOC argument gets flipped internally to give the more widely adopted convention for the SOC direction.

Returns

The evaluation of the derivative of the inverse model function.

5.12.2.10 inverse_d_dSOC_OCV_fit_function() `def ep_bolfi.utility.fitting_functions.inverse_d_dSOC_OCV_fit_function (`
`SOC,`
`* args,`
`z = 1.0,`
`T = 298.15,`
`inverted = True)`

The derivative of the inverse of [fitting_functions.OCV_fit_function](#).

Basically OCV'(SOC). Requires that the Δx entries are sorted by x. This corresponds to the parameters being sorted by decreasing E_0 .

Parameters

<i>E_OCV</i>	The SOC's for which the voltages shall be evaluated.
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Parameters

<i>args</i>	A list which length is dividable by three. These are the parameters E_0 , a and Δx from "A parametric OCV model" in the order (E_{0_0} , a_0 , Δx_0 , E_{0_1} , a_1 , $\Delta x_1...$).
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>inverted</i>	The default (False) uses the formulation from "A parametric OCV model". If True, the SOC argument gets flipped internally to give the more widely adopted convention for the SOC direction.

Returns

The evaluation of the derivative of the inverse model function.

5.12.2.11 inverse_OCV_fit_function() `def ep_bolfi.utility.fitting_functions.inverse_OCV_fit_function (`
 SOC,
 * *args*,
 z = 1.0,
 T = 298.15,
 inverted = True)

The inverse of [fitting_functions.OCV_fit_function](#).

Basically OCV(SOC). Requires that the Δx entries are sorted by x . This corresponds to the parameters being sorted by decreasing E_0 .

Parameters

<i>E_OCV</i>	The SOC's for which the voltages shall be evaluated.
<i>args</i>	A list which length is dividable by three. These are the parameters E_0 , a and Δx from "A parametric OCV model" in the order (E_{0_0} , a_0 , Δx_0 , E_{0_1} , a_1 , $\Delta x_1...$).
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>inverted</i>	The default (False) uses the formulation from "A parametric OCV model". If True, the SOC argument gets flipped internally to give the more widely adopted convention for the SOC direction.

Returns

The evaluation of the inverse model function.

5.12.2.12 laplace_transform() `def ep_bolfi.utility.fitting_functions.laplace_transform (`
 x,
 y,
 s)

Performs a basic laplace transformation.

Parameters

x	The independent variable.
y	The dependent variable.
s	The (possibly complex) frequencies for which to perform the transform.

Returns

The evaluation of the laplace transform at s .

5.12.2.13 OCV_fit_function() `def ep_bolfi.utility.fitting_functions.OCV_fit_function (`
`E_OCV,`
`* args,`
`z = 1.0,`
`T = 298.15,`
`individual = False,`
`fit_SOC_range = False,`
`rescale = False)`

The OCV model from "A parametric OCV model".

5.12.2.14 Reference C. R. Birkl, E. McTurk, M. R. Roberts, P. G. Bruce and D. A. Howey. "A Parametric Open Circuit Voltage Model for Lithium Ion Batteries". Journal of The Electrochemical Society, 162(12):A2271-A2280, 2015

Parameters

<i>E_OCV</i>	The voltages for which the SOC's shall be evaluated.
<i>args</i>	A list which length is dividable by three. These are the parameters E_0 , a and Δx from the paper referenced above in the order (E_{0_0} , a_0 , Δx_0 , E_{0_1} , a_1 , Δx_1 ...). If "fit_SOC_range" is True, two additional arguments are at the front (see there). The last Δx_j may be omitted to force $\sum \Delta x_j = 1$.
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>individual</i>	If True, the model function summands are not summed up.
<i>fit_SOC_range</i>	If True, this function takes two additional arguments at the start of "args" that may be used to adjust the data SOC range.
<i>rescale</i>	If True, the expected "args" now contain the slopes of the summands at their respective origins instead of a . Formula: $a / \text{slope} = -4 * (k_B / e) * T / (\Delta x * z)$.

Returns

The evaluation of the model function. This is the referenced fit function on a linearly transformed SOC range if "fit_SOC_range" is True. If "individual" is True, the individual summands in the model function get returned as a list.

5.12.2.15 `smooth_fit()` `def ep_bolfi.utility.fitting_functions.smooth_fit (`
`x,`
`y,`
`order = 3,`
`splits = None,`
`w = None,`
`s = None,`
`display = False,`
`derivatives = 0)`

Calculates a smoothed spline with derivatives.

Note: the "roots" function of a spline only works if it is cubic, i.e. of third order. Each "derivative" reduces the order by one.

Parameters

<i>x</i>	The independent variable.
<i>y</i>	The dependent variable ("plotted over x").
<i>order</i>	Interpolation order of the spline.
<i>splits</i>	Optional tuning parameter. A list of points between which splines will be fitted first. The returned spline then is a fit of these individual splines.
<i>w</i>	Optional list of weights. Works best if 1 / w approximates the standard deviation of the noise in y at each point. Defaults to SciPy default behaviour of <code>scipy.interpolate.UnivariateSpline</code> .
<i>s</i>	Optional tuning parameter. Higher values lead to coarser, but more smooth interpolations and vice versa. Defaults to SciPy default behaviour of <code>scipy.interpolate.UnivariateSpline</code> .
<i>display</i>	If set to True, the fit parameters of the spline will be printed to console. If possible, a monomial representation is printed.
<i>derivatives</i>	The derivatives of the spline to also include in the return. Default is 0, which gives the spline. 1 would give the spline, followed by its derivative. Can not be higher than spline order. Derivatives are only continuous when <code>derivatives < order</code> .

Returns

A smoothing spline in the form of `scipy.UnivariateSpline`.

5.12.2.16 `verbose_spline_parameterization()` `def ep_bolfi.utility.fitting_functions.verbose_spline_parameterization (`
`coeffs,`
`knots,`
`order,`
`format = 'python',`
`function_name = "OCV",`
`function_args = "SOC",`
`derivatives = 0,`
`spline_transformation = "",`
`verbose = False)`

Gives the monomic representation of a B-spline.

Parameters

<i>coeffs</i>	The B-spline coefficients as structured by <code>scipy.interpolate</code> .
<i>knots</i>	The B-spline knots as structured by <code>scipy.interpolate</code> .
<i>order</i>	The order of the B-spline.
<i>format</i>	Gives the file/language format for the function representation. Default is 'python'. The other choice is 'matlab'.
<i>function_name</i>	An optional name for the printed Python function.
<i>function_args</i>	An optional string for the arguments of the function.
<i>derivatives</i>	The derivatives of the spline to also include in the return. Default is 0, which gives the spline. 1 would give the spline, followed by its derivative. Can not be higher than spline order. Derivatives are only continuous when <code>derivatives < order</code> .
<i>spline_transformation</i>	Give a string if you want to include a function that gets applied to the whole spline, e.g. 'exp'.
<i>verbose</i>	Print information about the progress of the conversion.

Returns

A string that gives a Python function when "exec"-uted.

5.13 `ep_bolfi.utility.preprocessing` Namespace Reference

Contains frequently used workflows in dataset preprocessing.

Classes

- class `SubstitutionDict`
A dictionary with some automatic substitutions.

Functions

- def `fix_parameters` (`parameters_to_be_fixed`)
Returns a function which sets some parameters in advance.
- def `combine_parameters_to_try` (`parameters`, `parameters_to_try_dict`)
Give every combination as full parameter sets.
- def `calculate_means_and_standard_deviations` (`mean`, `covariance`, `free_parameters_names`, `transform_↔`
`parameters={}`, `bounds_in_standard_deviations=1`, `**kwargs`)
Calculate means and standard deviations.
- def `approximate_confidence_ellipsoid` (`parameters`, `free_parameters_names`, `covariance`, `mean=None`,
`transform_parameters={}`, `refinement=True`, `confidence=0.95`)
Approximate a confidence ellipsoid.
- def `capacity` (`parameters`, `electrode="positive"`)
Convenience function for calculating the capacity.
- def `calculate_SOC` (`timepoints`, `currents`, `initial_SOC=0`, `sign=1`, `capacity=1`)
Transforms applied current over time into SOC.
- def `calculate_both_SOC_from_OCV` (`parameters`, `negative_SOC_from_cell_SOC`, `positive_SOC_from_↔`
`cell_SOC`, `OCV`)
Calculates the SOC of both electrodes from their OCV.

- def [subtract_OCV_curve_from_cycles](#) (dataset, parameters, starting_SOC=None, starting_OCV=None, electrode="positive", current_sign=0, voltage_sign=0)
Removes the OCV curve from a cycling measurement.
- def [subtract_both_OCV_curves_from_cycles](#) (dataset, parameters, negative_SOC_from_cell_SOC, positive_SOC_from_cell_SOC, starting_SOC=None, starting_OCV=None)
Removes the OCV curve from a single cycle.
- def [laplace_transform](#) (x, y, s)
Performs a basic laplace transformation.
- def [find_occurrences](#) (sequence, value)
Gives indices in sequence where it is closest to value.
- def [OCV_from_CC_CV](#) (charge, cv, discharge, name, phases, eval_points=200, spline_SOC_range=(0.01, 0.99), spline_order=2, spline_smoothing=2e-3, spline_print=None, parameters_print=False)
Tries to extract the OCV curve from CC-CV cycling data.
- def [calculate_desired_voltage](#) (solution, t_eval, voltage_scale, overpotential, three_electrode=None, dimensionless_reference_electrode_location=0.5, parameters={})
- def [solve_all_parameter_combinations](#) (model, t_eval, parameters, parameters_to_try, submesh_types, var_pts, spatial_methods, full_factorial=True, **kwargs)
- def [prepare_parameter_combinations](#) (parameters, parameters_to_try, covariance, order_of_parameter_names, transform_parameters, confidence)
Calculates all permutations of the parameter boundaries.
- def [parallel_simulator_with_setup](#) (model, current_input, parameters, submesh_types, var_pts, spatial_methods, calc_esoh, inputs, t_eval, voltage_scale, overpotential, three_electrode, dimensionless_reference_electrode_location, kwargs)
- def [simulate_all_parameter_combinations](#) (model, current_input, submesh_types, var_pts, spatial_methods, parameters, parameters_to_try=None, covariance=None, order_of_parameter_names=None, additional_input_parameters=[], transform_parameters={}, confidence=0.95, full_factorial=True, calc_esoh=False, voltage_scale=1.0, overpotential=False, three_electrode=None, dimensionless_reference_electrode_location=0.5, **kwargs)

5.13.1 Detailed Description

Contains frequently used workflows in dataset preprocessing.

The functions herein are a collection of simple, but frequent, transformations of arrays of raw measurement data.

5.13.2 Function Documentation

5.13.2.1 [approximate_confidence_ellipsoid\(\)](#) `def ep_bolfi.utility.preprocessing.approximate_confidence_ellipsoid (`
`parameters,`
`free_parameters_names,`
`covariance,`
`mean = None,`
`transform_parameters = {},`
`refinement = True,`
`confidence = 0.95)`

Approximate a confidence ellipsoid.

Compatible with [SubstitutionDict](#), if "parameters" is one. The geometric approximation is a refinement of the polytope with nodes on the semiaxes of the confidence ellipsoid. The refinement step adds a node for each face, i.e., each sub-polytope with dimension smaller by 1. This node is centered on that face and projected onto the confidence ellipsoid.

Parameters

<i>parameters</i>	The base full parameter set as a dictionary.
<i>free_parameters_names</i>	The names of the parameters that are uncertain as a list. This parameter has to match the order of parameters in "covariance".
<i>covariance</i>	The covariance of the uncertain parameters as a two-dimensional numpy array.
<i>mean</i>	The mean of the uncertain parameters as a dictionary. If not set, the values from 'parameters' will be used.
<i>transform_parameters</i>	Optional transformations between the parameter space that is used for searching for optimal parameters and the model parameters. Any missing free parameter is not transformed. The values are 2-tuples. The first entry is a function taking the search space parameter and returning the model parameter. The second entry is the inverse function. For convenience, any value may also be one of the following: <ul style="list-style-type: none"> • 'none' => (identity, identity) • 'log' => (exp, log)
<i>confidence</i>	The confidence within the ellipsoid. Defaults to 0.95, i.e., the 95% confidence ellipsoid.
<i>refinement</i>	If False, only the nodes on the semiaxes get returned. If True, the nodes centered on the faces get returned as well.

Returns

A 2-tuple where the first item is the list of all parameter set combinations and the second the ellipsoid nodes only as a two-dimensional numpy array with each node in on row.

5.13.2.2 calculate_both_SOC_from_OCV() `def ep_bolfi.utility.preprocessing.calculate_both_SOC_from_OCV (`
parameters,
negative_SOC_from_cell_SOC,
positive_SOC_from_cell_SOC,
OCV)

Calculates the SOC of both electrodes from their OCV.

The SOC's are substituted in the given "parameters". The SOC of the cell as a whole gets returned in case it is needed.

Parameters

<i>parameters</i>	The parameters of the battery as used for the PyBaMM simulations (see models.standard_parameters).
<i>negative_SOC_from_cell_SOC</i>	A function that takes the SOC of the cell and returns the SOC of the negative electrode.
<i>positive_SOC_from_cell_SOC</i>	A function that takes the SOC of the cell and returns the SOC of the positive electrode.
<i>OCV</i>	The OCV for which the SOC's shall be calculated.

Returns

The SOC of the cell as a whole.

5.13.2.3 calculate_desired_voltage() `def ep_bolfi.utility.preprocessing.calculate_desired_voltage (`
solution,
t_eval,
voltage_scale,
overpotential,
three_electrode = None,
dimensionless_reference_electrode_location = 0.5,
parameters = {})

Parameters

<i>solution</i>	The pybamm.Solution object from which to calculate the voltage.
<i>t_eval</i>	The times at which to evaluate the "solution".
<i>voltage_scale</i>	The returned voltage gets divided by this value. For example, 1e-3 would produce a plot in [mV].
<i>overpotential</i>	If True, only the overpotential of "solutions" gets plotted. Otherwise, the cell voltage (OCV + overpotential) is plotted.
<i>three_electrode</i>	With None, does nothing (i.e., cell potentials are used). If set to either 'positive' or 'negative', instead of cell potentials, the base for the displayed voltage will be the potential of the 'positive' or 'negative' electrode against a reference electrode. For placement of said reference electrode, please refer to "dimensionless_reference_electrode_location".
<i>dimensionless_reference_electrode_location</i>	The location of the reference electrode, given as a scalar between 0 (placed at the point where negative electrode and separator meet) and 1 (placed at the point where positive electrode and separator meet). Defaults to 0.5 (in the middle).
<i>parameters</i>	The parameter dictionary that was used for the simulation. Only needed for a three-electrode output.

Returns

The array of the specified voltages over time.

5.13.2.4 calculate_means_and_standard_deviations() `def ep_bolfi.utility.preprocessing.calculate_means_and_↵`
standard_deviations (
mean,
covariance,
free_parameters_names,
transform_parameters = {},
bounds_in_standard_deviations = 1,
*** kwargs)*

Calculate means and standard deviations.

Please note that standard deviations translate differently into confidence regions in different dimensions. For the confidence region, use "approximate_confidence_ellipsoid".

Parameters

<i>mean</i>	The mean of the uncertain parameters as a dictionary.
<i>covariance</i>	The covariance of the uncertain parameters as a two-dimensional numpy array.
<i>free_parameters_names</i>	The names of the parameters that are uncertain as a list. This parameter maps the order of parameters in "covariance".
<i>transform_parameters</i>	Optional transformations between the parameter space that is used for searching for optimal parameters and the model parameters. Any missing free parameter is not transformed. The values are 2-tuples. The first entry is a function taking the search space parameter and returning the model parameter. The second entry is the inverse function. For convenience, any value may also be one of the following: <ul style="list-style-type: none"> • 'none' => (identity, identity) • 'log' => (exp, log)
<i>bounds_in_standard_deviations</i>	Sets how many standard deviations in each direction the returned error bounds are. These are first applied and then transformed.
<i>**kwargs</i>	Keyword arguments for <code>scipy.integrate.quad</code> , which is used to numerically calculate mean and variance.

Returns

A 3-tuple with three dictionaries. Their keys are the free parameters' names as keys and their values are those parameters' means, standard deviations, and error bounds.

5.13.2.5 calculate_SOC() `def ep_bolfi.utility.preprocessing.calculate_SOC (`
timepoints,
currents,
initial_SOC = 0,
sign = 1,
capacity = 1)

Transforms applied current over time into SOC.

Parameters

<i>timepoints</i>	Array of the timepoint segments.
<i>currents</i>	Array of the current segments.
<i>initial_SOC</i>	The SOC value to start accumulating from.
<i>sign</i>	The value by which to multiply the current.
<i>capacity</i>	A scaling by which to convert from C to dimensionless SOC.

Returns

An array of the same shape describing SOC in C.

5.13.2.6 capacity() `def ep_bolfi.utility.preprocessing.capacity (`
 parameters,
 electrode = "positive")

Convenience function for calculating the capacity.

Parameters

<i>parameters</i>	A parameter file as defined by models.standard_parameters .
<i>electrode</i>	The prefix of the electrode to use for capacity calculation. Change to "negative" to use the one with the lower OCP.

Returns

The capacity of the parameterized battery in C.

5.13.2.7 combine_parameters_to_try() `def ep_bolfi.utility.preprocessing.combine_parameters_to_try (`
 parameters,
 parameters_to_try_dict)

Give every combination as full parameter sets.

Compatible with [SubstitutionDict](#), if "parameters" is one.

Parameters

<i>parameters</i>	The base full parameter set as a dictionary.
<i>parameters_to_try_dict</i>	The keys of this dictionary correspond to the "parameters" keys where different values are to be inserted. These are given by the tuples which are the values of this dictionary.

Returns

A 2-tuple where the first item is the list of all parameter set combinations and the second the list of the combinations only.

5.13.2.8 find_occurrences() `def ep_bolfi.utility.preprocessing.find_occurrences (`
 sequence,
 value)

Gives indices in sequence where it is closest to value.

Parameters

<i>sequence</i>	A list that represents a differentiable function.
<i>value</i>	The value that is searched for in "sequence". Also, crossings of consecutive values in "sequence" with "value" are searched for.

Returns

A list of indices in "sequence" in ascending order where "value" or a close match for "value" was found.

5.13.2.9 fix_parameters() `def ep_bolfi.utility.preprocessing.fix_parameters (`
 parameters_to_be_fixed)

Returns a function which sets some parameters in advance.

Parameters

<i>parameters_to_be_fixed</i>	These parameters will at least be a part of the dictionary that the returned function returns.
-------------------------------	--

Returns

The function which adds additional parameters to a dictionary or replaces existing parameters with the new ones.

5.13.2.10 laplace_transform() `def ep_bolfi.utility.preprocessing.laplace_transform (`
 x,
 y,
 s)

Performs a basic laplace transformation.

Parameters

<i>x</i>	The independent variable.
<i>y</i>	The dependent variable.
<i>s</i>	The (possibly complex) frequencies for which to perform the transform.

Returns

The evaluation of the laplace transform at *s*.

5.13.2.11 OCV_from_CC_CV() `def ep_bolfi.utility.preprocessing.OCV_from_CC_CV (`
 charge,
 cv,
 discharge,
 name,
 phases,
 eval_points = 200,
 spline_SOC_range = (0.01, 0.99),

```
spline_order = 2,
spline_smoothing = 2e-3,
spline_print = None,
parameters_print = False)
```

Tries to extract the OCV curve from CC-CV cycling data.

Parameters

<i>charge</i>	A Cycling_Information object containing the constant charge cycle(s). If more than one CC-CV-cycle shall be analyzed, please make sure that the order of this, cv and discharge align.
<i>cv</i>	A Cycling_Information object containing the constant voltage part between charge and discharge cycle(s).
<i>discharge</i>	A Cycling_Information object containing the constant discharge cycle(s). These occur after each cv cycle.
<i>name</i>	Name of the material for which the CC-CV-cycling was measured.
<i>phases</i>	Number of phases in the fitting_functions.OCV_fit_function as an int. The higher it is, the more (over-)fitted the model becomes.
<i>eval_points</i>	The number of points for plotting of the OCV curves.
<i>spline_SOC_range</i>	2-tuple giving the SOC range in which the inverted fitting_functions.OCV_fit_function will be interpolated by a smoothing spline. Outside of this range the spline is used for extrapolation. Use this to fit the SOC range of interest more precisely, since a fit of the whole range usually fails due to the singularities at SOC 0 and 1. Please note that this range considers the 0-1-range in which the given SOC lies and not the linear transformation of it from the fitting process.
<i>spline_order</i>	Order of this smoothing spline. If it is set to 0, only the fitting_functions.OCV_fit_function is calculated and plotted.
<i>spline_smoothing</i>	Smoothing factor for this smoothing spline. Default: 2e-3. Lower numbers give more precision, while higher numbers give a simpler spline that smoothes over steep steps in the fitted OCV curve.
<i>spline_print</i>	If set to either 'python' or 'matlab', a string representation of the smoothing spline is printed in the respective format.
<i>parameters_print</i>	Set to True if the fit parameters should be printed to console.

Returns

A 8-tuple consisting of the following: 0: OCV_fits The fitted OCV curve parameters for each CC-CV cycle as returned by [fitting_functions.fit_OCV](#). 1: I_mean The currents assigned to each CC-CV cycle (without CV). 2: C_charge The moved capacities during the charge segment(s). This is a list of the same length as charge, cv or discharge. 3: U_charge The voltages during the charge segment(s). Length: same. 4: C_discharge The moved capacities during the discharge segment(s). Length: same. 5: U_discharge The voltages during the discharge segment(s). Length: same. 6: C_evals Structurally the same as C_charge or C_discharge, this contains the moved capacities that were assigned to the mean voltages of charge and discharge cycle(s). 7: U_means The mean voltages of each charge and discharge cycle.

5.13.2.12 parallel_simulator_with_setup() `def ep_bolfi.utility.preprocessing.parallel_simulator_with_setup (`
model,
current_input,
parameters,

```

    submesh_types,
    var_pts,
    spatial_methods,
    calc_esoh,
    inputs,
    t_eval,
    voltage_scale,
    overpotential,
    three_electrode,
    dimensionless_reference_electrode_location,
    kwargs )

```

5.13.2.13 prepare_parameter_combinations() `def ep_bolfi.utility.preprocessing.prepare_parameter_combinations (`
parameters,
parameters_to_try,
covariance,
order_of_parameter_names,
transform_parameters,
confidence)

Calculates all permutations of the parameter boundaries.

Parameters

<i>parameters</i>	The model parameters as a dictionary.
<i>parameters_to_try</i>	A dictionary with the names of the model parameters as keys and lists of the values that are to be tried out for them as values. Mutually exclusive to "covariance".
<i>covariance</i>	A covariance matrix describing an estimation result of model parameters. Will be used to calculate parameters to try that together approximate the confidence ellipsoid. This confidence ellipsoid will be centered on "parameters". Mutually exclusive to "parameters_to_try".
<i>order_of_parameter_names</i>	A list of names from "parameters" that correspond to the order these parameters appear in the rows and columns of "covariance". Only needed when "covariance" is set.
<i>transform_parameters</i>	Optional transformations between the parameter space that is used for searching for optimal parameters and the model parameters. Any missing free parameter is not transformed. The values are 2-tuples. The first entry is a function taking the search space parameter and returning the model parameter. The second entry is the inverse function. For convenience, any value may also be one of the following: <ul style="list-style-type: none"> • 'none' => (identity, identity) • 'log' => (exp, log)
<i>confidence</i>	The confidence within the ellipsoid. Defaults to 0.95, i.e., the 95% confidence ellipsoid.

Returns

A 2-tuple with the individual parameter variations and then all permutations of them.

5.13.2.14 simulate_all_parameter_combinations() `def ep_bolfi.utility.preprocessing.simulate_all_parameter_↵`
 combinations (
 model,
 current_input,
 submesh_types,
 var_pts,
 spatial_methods,
 parameters,
 parameters_to_try = None,
 covariance = None,
 order_of_parameter_names = None,
 additional_input_parameters = [],
 transform_parameters = {},
 confidence = 0.95,
 full_factorial = True,
 calc_esoh = False,
 voltage_scale = 1.0,
 overpotential = False,
 three_electrode = None,
 dimensionless_reference_electrode_location = 0.5,
 ** *kwargs*)

Parameters

<i>model</i>	The PyBaMM battery model that is to be solved.
<i>current_input</i>	The list of battery operation conditions. See <code>pybamm.Simulation</code> .
<i>submesh_types</i>	The submeshes for discretization. See <code>solversetup.spectral_mesh_pts_and_method</code> .
<i>var_pts</i>	The number of discretization points. See <code>solversetup.spectral_mesh_pts_and_method</code> .
<i>spatial_methods</i>	The spatial methods for discretization. See <code>solversetup.spectral_mesh_pts_and_method</code> .
<i>parameters</i>	The model parameters as a dictionary.
<i>parameters_to_try</i>	A dictionary with the names of the model parameters as keys and lists of the values that are to be tried out for them as values. Mutually exclusive to "covariance".
<i>covariance</i>	A covariance matrix describing an estimation result of model parameters. Will be used to calculate parameters to try that together approximate the confidence ellipsoid. This confidence ellipsoid will be centered on "parameters". Mutually exclusive to "parameters_to_try".
<i>order_of_parameter_names</i>	A list of names from "parameters" that correspond to the order these parameters appear in the rows and columns of "covariance". Only needed when "covariance" is set.
<i>additional_input_parameters</i>	A list of the parameter names that are changed by any of the variable parameters, if "parameters" is a SubstitutionDict .
<i>transform_parameters</i>	Optional transformations between the parameter space that is used for searching for optimal parameters and the model parameters. Any missing free parameter is not transformed. The values are 2-tuples. The first entry is a function taking the search space parameter and returning the model parameter. The second entry is the inverse function. For convenience, any value may also be one of the following: <ul style="list-style-type: none"> • 'none' => (identity, identity) • 'log' => (exp, log)

Parameters

<i>confidence</i>	The confidence within the ellipsoid. Defaults to 0.95, i.e., the 95% confidence ellipsoid.
<i>full_factorial</i>	When "parameters_to_try" is set: <ul style="list-style-type: none"> • If True, all parameter combinations are tried out. If False, only each parameter is varied with the others staying fixed. When "covariance" is set: • If False, only the points on the semiaxes of the confidence ellipsoid constitute the parameters to try. If True, the centres of the faces of the polytope of these points get added to the parameters to try, projected onto the surface of the confidence ellipsoid.
<i>calc_esoh</i>	Passed on to pybamm.Simulator, see there.
<i>kwargs</i>	The optional parameters for solversetup.solver_setup. See there.

Returns

A 2-tuple with the model solution for parameters as first entry. The second entry mimics parameters_to_try with each entry in their lists replaced by the model solution for the corresponding parameter substitution. The second entry has one additional key "all parameters", where all parameters_to_try combinations are the value.

5.13.2.15 solve_all_parameter_combinations() `def ep_bolfi.utility.preprocessing.solve_all_parameter_combinations (`
`model,`
`t_eval,`
`parameters,`
`parameters_to_try,`
`submesh_types,`
`var_pts,`
`spatial_methods,`
`full_factorial = True,`
`** kwargs)`

Parameters

<i>model</i>	The PyBaMM battery model that is to be solved.
<i>t_eval</i>	The timepoints in s at which this model is to be solved.
<i>parameters</i>	The model parameters as a dictionary.
<i>parameters_to_try</i>	A dictionary with the names of the model parameters as keys and lists of the values that are to be tried out for them as values.
<i>submesh_types</i>	The submeshes for discretization. See solversetup.spectral_mesh_pts_and_method.
<i>var_pts</i>	The number of discretization points. See solversetup.spectral_mesh_pts_and_method.
<i>spatial_methods</i>	The spatial methods for discretization. See solversetup.spectral_mesh_pts_and_method.
<i>full_factorial</i>	If True, all parameter combinations are tried out. If False, only each parameter is varied with the others staying fixed.
<i>kwargs</i>	The optional parameters for solversetup.solver_setup. See there.

Returns

A 2-tuple with the model solution for parameters as first entry. The second entry mimics `parameters_to_try` with each entry in their lists replaced by the model solution for the corresponding parameter substitution. The second entry has one additional key "all parameters", where all `parameters_to_try` combinations are the value.

5.13.2.16 subtract_both_OCV_curves_from_cycles() `def ep_bolfi.utility.preprocessing.subtract_both_OCV_curves←↵
_from_cycles (`
`dataset,`
`parameters,`
`negative_SOC_from_cell_SOC,`
`positive_SOC_from_cell_SOC,`
`starting_SOC = None,`
`starting_OCV = None)`

Removes the OCV curve from a single cycle.

Parameters

<i>dataset</i>	A Cycling_Information object of the measurement.
<i>parameters</i>	The parameters of the battery as used for the PyBaMM simulations (see models.standard_parameters).
<i>negative_SOC_from_cell_SOC</i>	A function that takes the SOC of the cell and returns the SOC of the negative electrode.
<i>positive_SOC_from_cell_SOC</i>	A function that takes the SOC of the cell and returns the SOC of the positive electrode.
<i>starting_SOC</i>	The SOC at the beginning of the measurement. If not given, the OCV curves will be inverted to determine the initial SOC.
<i>starting_OCV</i>	The OCV at the beginning of the measurement. If not given, the first entry of voltages is used for this.

Returns

2-tuple. First entry are the voltages minus the OCV as estimated for each data point. These are structured in exactly the same way as in the "dataset". Second entry are the electrode SOC's as counted in the data.

5.13.2.17 subtract_OCV_curve_from_cycles() `def ep_bolfi.utility.preprocessing.subtract_OCV_curve_from_cycles (`
`dataset,`
`parameters,`
`starting_SOC = None,`
`starting_OCV = None,`
`electrode = "positive",`
`current_sign = 0,`
`voltage_sign = 0)`

Removes the OCV curve from a cycling measurement.

Parameters

<i>dataset</i>	A Cycling_Information object of the measurement.
<i>parameters</i>	The parameters of the battery as used for the PyBaMM simulations (see models.standard_parameters).
<i>starting_SOC</i>	The SOC at the beginning of the measurement. If not given, the OCV curve will be inverted to determine the initial SOC.
<i>starting_OCV</i>	The OCV at the beginning of the measurement. If not given and starting_SOC is also not given, the first entry of voltages is used for this. If not given, but starting_SOC is, the OCP function will be evaluated at starting_SOC to get the OCV.
<i>electrode</i>	"positive" (default) or "negative" for current sign correction and capacity calculation. "positive" adds SOC with positive current and vice versa. The sign corrections can be overwritten with '*_sign'.
<i>current_sign</i>	1 adds SOC, -1 subtracts it, 0 follows the default behaviour above.
<i>voltage_sign</i>	1 subtracts the OCP, -1 adds it, 0 follows the default behaviour above.

Returns

2-tuple. First entry are the voltages minus the OCV as estimated for each data point. These are structured in exactly the same way as in the "dataset". Second entry are the electrode SOC's as counted in the data.

5.14 ep_bolfi.utility.visualization Namespace Reference

Various helper and plotting functions for common data visualizations.

Functions

- def [update_limits](#) (ax, xmin=float('inf'), xmax=-float('inf'), ymin=float('inf'), ymax=-float('inf'))
Convenience function for adjusting the view.
- def [set_fontsize](#) (ax, title=12, xaxis=12, yaxis=12, xticks=12, yticks=12, legend=12)
Convenience function for fontsize changes.
- def [update_legend](#) (ax, additional_handles=[], additional_labels=[], additional_handler_map={})
Makes sure that all items remain and all items show up.
- def [push_apart_text](#) (fig, ax, text_objects, lock_xaxis=False, temp_path="/temp_render.png")
Push apart overlapping texts until no overlaps remain.
- def [make_segments](#) (x, y)
Create a list of line segments from x and y coordinates.
- def [colorline](#) (x, y, z=None, cmap=plt.get_cmap('viridis'), norm=matplotlib.colors.Normalize(0, 1), linewidth=1, linestyle='-', alpha=1.0)
Generates a colored line using LineCollection.
- def [nyquist_plot](#) (fig, ax, ω , Z, cmap=plt.get_cmap('tab20b'), ls='-', lw=3, title_text="Impedance Measurement", legend_text="impedance", colorbar_label="Frequency / Hz", add_frequency_colorbar=True, equal_aspect=True)
Plot an impedance measurement.
- def [bode_plot](#) (fig, ax_real, ax_imag, ω , Z, cmap=plt.get_cmap('tab20b'), ls_real='-', ls_imag='-', lw=3, title_text="Impedance Measurement", legend_text="impedance")
Plot an impedance measurement.
- def [plot_comparison](#) (ax, solutions, errorbars, experiment, solution_visualization=[], t_eval=None, title="", xlabel="", ylabel="", feature_visualizer=lambda *args:[], feature_fontsize=12, interactive_plot=False, output_variables=None, voltage_scale=1.0, use_cycles=False, overpotential=False, three_electrode=None, dimensionless_reference_electrode_location=0.5, parameters=None)

Tool for comparing simulation<->experiment with features.

- `def cc_cv_visualization` (fig, ax, dataset, max_number_of_clusters=4, cmap=plt.get_cmap('tab20c'), check_location=[0.1, 0.7, 0.2, 0.225])

Automatically labels and displays a CC-CV dataset.

- `def plot_OCV_from_CC_CV` (ax_ICA_meas, ax_ICA_mean, ax_OCV_meas, ax_OCV_mean, charge, cv, discharge, name, phases, eval_points=200, spline_SOC_range=(0.01, 0.99), spline_order=2, spline_smoothing=2e-3, spline_print=None, parameters_print=False)

Visualizes the OCV_fitting.OCV_from_CC_CV output.

- `def plot_ICA` (ax, SOC, OCV, name, spline_order=2, spline_smoothing=2e-3, sign=1)

Show the derivative of charge by voltage.

- `def plot_measurement` (fig, ax, dataset, title, cmap=plt.get_cmap('tab20c'), plot_current=True)

Plots current and voltage curves in one diagram.

- `def fit_and_plot_OCV` (ax, SOC, OCV, name, phases, SOC_range_bounds=(0.2, 0.8), SOC_range_limits=(0.0, 1.0), z=1.0, T=298.15, fit=None, eval_SOC=[0, 1], eval_points=200, spline_SOC_range=(0.01, 0.99), spline_order=2, spline_print=None, parameters_print=False, inverted=True, info_accuracy=True, normalized_xaxis=False, distance_order=2, weights=None, initial_parameters=None, minimize_options=None)

Fits an SOC(OCV)-model and an OCV(SOC)-evaluable spline.

- `def visualize_correlation` (fig, ax, correlation, names=None, title=None, cmap=plt.get_cmap('BrBG'), entry_color='w')

Produces a heatmap of a correlation matrix.

5.14.1 Detailed Description

Various helper and plotting functions for common data visualizations.

5.14.2 Function Documentation

5.14.2.1 `bode_plot()`

```
def ep_bolfi.utility.visualization.bode_plot (
    fig,
    ax_real,
    ax_imag,
     $\omega$ ,
    Z,
    cmap = plt.get_cmap('tab20b'),
    ls_real = '-',
    ls_imag = '-.',
    lw = 3,
    title_text = "Impedance Measurement",
    legend_text = "impedance")
```

Plot an impedance measurement.

Parameters

<i>fig</i>	The matplotlib.Figure for plotting
<i>ax</i>	The matplotlib.Axes for plotting.
ω	The frequencies at which the impedance was measured. May be a list of lists for multiple measurements.

Parameters

<i>Z</i>	The impedances that were measured at those frequencies. May be a list of lists for multiple measurements.
<i>cmap</i>	The colormap that is used to differentiate multiple impedances.
<i>ls_real</i>	The linestyle of the plot of the real part of the impedance.
<i>ls_imag</i>	The linestyle of the plot of the imaginary part of the impedance.
<i>lw</i>	The linewidth of the plot.
<i>title_text</i>	The text for the title.
<i>legend_text</i>	The text for the legend. May be a list for multiple measurements.

5.14.2.2 cc_cv_visualization() def ep_bolfi.utility.visualization.cc_cv_visualization (
fig,
ax,
dataset,
max_number_of_clusters = 4,
cmap = plt.get_cmap('tab20c'),
check_location = [0.1, 0.7, 0.2, 0.225])

Automatically labels and displays a CC-CV dataset.

A checkbox list gets added for browsing through the labels.

Parameters

<i>fig</i>	The Figure where the check boxes shall be drawn.
<i>ax</i>	The Axes where the measurements shall be drawn.
<i>dataset</i>	An instance of Cycling_Information. Please refer to utility.dataset_formatting for further information.
<i>max_number_of_clusters</i>	The maximum number of different labels that shall be tried in the automatic labelling of the dataset.
<i>cmap</i>	The Colormap that is used for colorcoding the cycles.
<i>check_location</i>	The (x,y)-coordinates (first two entries) and the (width,height) (last two entries) of the checkbox list canvas.

Returns

The CheckButtons instance. The only thing that must be done with this is to keep it in memory. Otherwise, it gets garbage collected ("weak reference") and the checkboxes don't work.

5.14.2.3 colorline() def ep_bolfi.utility.visualization.colorline (
x,
y,
z = None,
cmap = plt.get_cmap('viridis'),
norm = matplotlib.colors.Normalize(0, 1),

```
linewidth = 1,
linestyle = '-.',
alpha = 1.0)
```

Generates a colored line using LineCollection.

<http://nbviewer.ipython.org/github/dpsanders/matplotlib-examples/blob/master/colorline.ipynb> http://matplotlib.org/examples/pylab_examples/multicolored_line.html

Parameters

<i>x</i>	The independent variable.
<i>y</i>	The dependent variable.
<i>z</i>	Specify colors.
<i>cmap</i>	Specify a colormap for colors.
<i>norm</i>	Specify a normalization for mapping <i>z</i> to the colormap. Example: <code>matplotlib.colors.LogNorm(10**(-2), 10**4)</code> .
<i>linewidth</i>	The linewidth of the generated LineCollection.
<i>linestyle</i>	The linestyle of the generated LineCollection. If the individual lines in there are too short, its effect might not be visible.
<i>alpha</i>	The transparency of the generated LineCollection.

Returns

A `matplotlib.collections.LineCollection` object "lc". It can be plotted by a Matplotlib axis *ax* with "`ax.add_collection(lc)`".

5.14.2.4 **fit_and_plot_OCV()**

```
def ep_bolfi.utility.visualization.fit_and_plot_OCV (
    ax,
    SOC,
    OCV,
    name,
    phases,
    SOC_range_bounds = (0.2, 0.8),
    SOC_range_limits = (0.0, 1.0),
    z = 1.0,
    T = 298.15,
    fit = None,
    eval_SOC = [0, 1],
    eval_points = 200,
    spline_SOC_range = (0.01, 0.99),
    spline_order = 2,
    spline_print = None,
    parameters_print = False,
    inverted = True,
    info_accuracy = True,
    normalized_xaxis = False,
    distance_order = 2,
    weights = None,
    initial_parameters = None,
    minimize_options = None)
```

Fits an SOC(OCV)-model and an OCV(SOC)-evaluable spline.

5.14.3 Exemplary fit parameters:

5.14.4 Fit parameters of a graphite anode.

```
E_0_g = np.array([0.35973, 0.17454, 0.12454, 0.081957])
yUeminus1_g = np.array([-0.33144, 8.9434e-3, 7.2404e-2, 6.7789e-2])
a_g = a_fit(yUeminus1_g)
Δx_g = np.array([8.041e-2, 0.23299, 0.29691, 0.39381])#0.22887
graphite = [p[i] for i in range(4) for p in [E_0_g, a_g, Δx_g]]
```

5.14.5 Fit parameters of a NMC-622 cathode.

```
E_0_NMC = np.array([4.2818, 3.9632, 3.9118, 3.6788])
yUeminus1_NMC = np.array([-0.22022, -0.083146, 0.070787, -0.11461])
a_NMC = a_fit(yUeminus1_NMC)
Δx_NMC = np.array([0.38646, 0.28229, 0.15104, 0.26562])#0.30105
NMC = [p[i] for i in range(4) for p in [E_0_NMC, a_NMC, Δx_NMC]]
```

Parameters

<i>ax</i>	The matplotlib.Axes instance for plotting.
<i>SOC</i>	Presumed SOC points of the OCV measurement. They only need to be precise in respect to relative capacity between measurements. The SOC endpoints of the measurement will be fitted using the fitting_functions.OCV_fit_function . Type: list or np.array.
<i>OCV</i>	OCV measurements as a list or np.array.
<i>name</i>	Name of the material for which the OCV curve was measured.
<i>phases</i>	Number of phases in the fitting_functions.OCV_fit_function as an int. The higher it is, the more (over-)fitted the model becomes.
<i>SOC_range_bounds</i>	Optional hard upper and lower bounds for the SOC correction from the left and the right side, respectively, as a 2-tuple. Use it as a limiting guess for the actual SOC range represented in the measurement. Has to be inside (0.0, 1.0). Set to (0.0, 1.0) to effectively disable SOC range estimation.
<i>SOC_range_limits</i>	Optional hard lower and upper bounds for the SOC correction from the left and the right side, respectively, as a 2-tuple. Use it if you know that your OCV data is incomplete and by how much. Has to be inside (0.0, 1.0). Set to (0.0, 1.0) to allow the SOC range estimation to assign datapoints to the asymptotes.
<i>z</i>	The charge number of the electrode interface reaction.
<i>T</i>	The temperature of the electrode.
<i>fit</i>	May provide the fit parameters if they are already known.
<i>eval_SOC</i>	Denotes the minimum and maximum SOC to plot the OCV curves at.
<i>eval_points</i>	The number of points for plotting of the OCV curves.
<i>spline_SOC_range</i>	2-tuple giving the SOC range in which the inverted fitting_functions.OCV_fit_function will be interpolated by a smoothing spline. Outside of this range the spline is used for extrapolation. Use this to fit the SOC range of interest more precisely, since a fit of the whole range usually fails due to the singularities at SOC 0 and 1. Please note that this range considers the 0-1-range in which the given SOC lies and not the linear transformation of it from the fitting process.
<i>spline_order</i>	Order of this smoothing spline. If it is set to 0, only the fitting_functions.OCV_fit_function is calculated and plotted.
<i>spline_print</i>	If set to either 'python' or 'matlab', a string representation of the smoothing spline is printed in the respective format.
<i>parameters_print</i>	Set to True if the fit parameters should be printed to console.
<i>inverted</i>	If True (default), the widely adopted SOC convention is assumed. If False, the formulation of "A parametric OCV model" is used.

Parameters

<i>info_accuracy</i>	If True, some measures of fit accuracy are displayed in the figure legend: RMSE (root mean square error), MAE (mean absolute error) and ME (maximum error).
<i>normalized_xaxis</i>	If True, the x-axis gets rescaled to [0,1], where {0,1} matches the asymptotes of the OCV fit function.
<i>distance_order</i>	The argument passed to the <code>numpy.linalg.norm</code> of the vector of distances between OCV data and OCV model. Default is 2, i.e., the Euclidean norm. 1 sets it to absolute distance.
<i>weights</i>	Optional weights to apply to the vector of the distances between OCV data and OCV model. Defaults to equal weights.
<i>initial_parameters</i>	Optional initial guess for the model parameters. If left as-is, this will be automatically gleaned from the data. Use only if you have another fit to data of the same electrode material.
<i>minimize_options</i>	Dictionary that gets passed to <code>scipy.optimize.minimize</code> with the method 'trust-constr'. See <code>scipy.optimize.show_options</code> with the arguments 'minimize' and 'trust-constr' for details.

5.14.5.1 make_segments() `def ep_bolfi.utility.visualization.make_segments (`
`x,`
`y)`

Create a list of line segments from x and y coordinates.

Parameters

<i>x</i>	The independent variable.
<i>y</i>	The dependent variable.

Returns

An array of the form `numlines x (points per line) times 2 (x and y) array`. This is the correct format for `LineCollection`.

5.14.5.2 nyquist_plot() `def ep_bolfi.utility.visualization.nyquist_plot (`
`fig,`
`ax,`
`ω ,`
`Z,`
`cmap = plt.get_cmap('tab20b'),`
`ls = '-',`
`lw = 3,`
`title_text = "Impedance Measurement",`
`legend_text = "impedance",`
`colorbar_label = "Frequency / Hz",`
`add_frequency_colorbar = True,`
`equal_aspect = True)`

Plot an impedance measurement.

Parameters

<i>fig</i>	The matplotlib.Figure for plotting
<i>ax</i>	The matplotlib.Axes for plotting.
ω	The frequencies at which the impedance was measured. May be a list of lists for multiple measurements.
<i>Z</i>	The impedances that were measured at those frequencies. May be a list of lists for multiple measurements.
<i>cmap</i>	The colormap that is used to visualize the frequencies.
<i>ls</i>	The linestyle of the plot.
<i>lw</i>	The linewidth of the plot.
<i>title_text</i>	The text for the title.
<i>legend_text</i>	The text for the legend. May be a list for multiple measurements.
<i>colorbar_label</i>	The label that is displayed next to the colorbar.
<i>add_frequency_colorbar</i>	Set to False if 'fig' was already decorated with a colorbar.
<i>equal_aspect</i>	Set to False in case the impedance is too vertical or horizontal.

Returns

A list of the LineCollection objects of the impedance plots.

5.14.5.3 plot_comparison() `def ep_bolfi.utility.visualization.plot_comparison (`

```

    ax,
    solutions,
    errorbars,
    experiment,
    solution_visualization = [],
    t_eval = None,
    title = "",
    xlabel = "",
    ylabel = "",
    feature_visualizer = lambda *args: [],
    feature_fontsize = 12,
    interactive_plot = False,
    output_variables = None,
    voltage_scale = 1.0,
    use_cycles = False,
    overpotential = False,
    three_electrode = None,
    dimensionless_reference_electrode_location = 0.5,
    parameters = None )

```

Tool for comparing simulation<->experiment with features.

First, a pybamm.QuickPlot shows the contents of "solutions". Then, a plot for feature visualization is generated.

Parameters

<i>ax</i>	The Axes onto which the comparison shall be plotted.
-----------	--

Parameters

<i>solutions</i>	A dictionary of pybamm.Solution objects. The key goes into the figure legend and the value gets plotted as a line.
<i>errorbars</i>	A dictionary of lists of either pybamm.Solution objects or lists of the desired variable at 't_eval' timepoints. The key goes into the figure legend and the values get plotted as a shaded area between the minimum and maximum.
<i>experiment</i>	A list/tuple of at least length 2. The first two entries are the data timepoints in s and voltages in V. The entries after that are only relevant as optional arguments to "feature_visualizer".
<i>solution_visualization</i>	This list/tuple is passed on to "feature_visualizer" in place of the additional entries of "experiment" for the visualization of the simulated features.
<i>t_eval</i>	The timepoints at which the "solutions" and "errorbars" shall be evaluated in s. If None are given, the timepoints of the solutions will be chosen.
<i>title</i>	The optional title of the feature visualization plot.
<i>xlabel</i>	The optional label of the x-axis there. Please note that the time will be given in h.
<i>ylabel</i>	The optional label of the y-axis there.
<i>feature_visualizer</i>	This is an optional function that takes "experiment" and returns a list of 2- or 3-tuples. The first two entries in the tuples are x- and y-data to be plotted alongside the other curves. The third entry is a string that is plotted at the respective (x[0], y[0])-coordinates.
<i>interactive_plot</i>	Choose whether or not a browsable overview of the solution components shall be shown. Please note that this disrupts the execution of this function until that plot is closed, since it is plotted in a new figure rather than in ax.
<i>output_variables</i>	The variables of "solutions" that are to be plotted. When None are specified, some default variables get plotted. The full list of possible variables to plot are returned by PyBaMM models from their get_fundamental_variables and get_coupled_variables functions. Enter the keys from that as strings in a list here.
<i>voltage_scale</i>	The plotted voltage gets divided by this value. For example, 1e-3 would produce a plot in [mV]. The voltage given to the "feature_visualizer" is not affected.
<i>use_cycles</i>	If True, the .cycles property of the "solutions" is used for the "feature_visualizer". Plotting is not affected.
<i>overpotential</i>	If True, only the overpotential of "solutions" gets plotted. Otherwise, the cell voltage (OCV + overpotential) is plotted.
<i>three_electrode</i>	By default, does nothing (i.e., cell potentials are used). If set to either 'positive' or 'negative', instead of cell potentials, the base for the displayed voltage will be the potential of the 'positive' or 'negative' electrode against a reference electrode. For placement of said reference electrode, please refer to "dimensionless_reference_electrode_location".
<i>dimensionless_reference_electrode_location</i>	The location of the reference electrode, given as a scalar between 0 (placed at the point where negative electrode and separator meet) and 1 (placed at the point where positive electrode and separator meet). Defaults to 0.5 (in the middle).
<i>parameters</i>	The parameter dictionary that was used for the simulation. Only needed for a three-electrode output.

Returns

The text objects that were generated according to "feature_visualizer".

5.14.5.4 plot_ICA() `def ep_bolfi.utility.visualization.plot_ICA (`
`ax,`
`SOC,`
`OCV,`
`name,`
`spline_order = 2,`
`spline_smoothing = 2e-3,`
`sign = 1)`

Show the derivative of charge by voltage.

Parameters

<i>ax</i>	The matplotlib.Axes instance for plotting.
<i>SOC</i>	Presumed SOC points of the OCV measurement. They only need to be precise in respect to relative capacity between measurements.
<i>OCV</i>	OCV measurements as a list or np.array, matching SOC.
<i>name</i>	Name of the material for which the OCV curve was measured.
<i>spline_order</i>	Order of the smoothing spline used for derivation. Default: 2.
<i>spline_smoothing</i>	Smoothing factor for this smoothing spline. Default: 2e-3. Lower numbers give more precision, while higher numbers give a simpler spline that smoothes over steep steps in the fitted OCV curve.
<i>sign</i>	Put -1 if the ICA comes out negative. Default: 1.

5.14.5.5 plot_measurement() `def ep_bolfi.utility.visualization.plot_measurement (`
`fig,`
`ax,`
`dataset,`
`title,`
`cmap = plt.get_cmap('tab20c'),`
`plot_current = True)`

Plots current and voltage curves in one diagram.

Please don't use "fig.tight_layout()" with this, as it very well might mess up the placement of the colorbar and the second y-axis. Rather, use "plt.subplots(..., constrained_layout=True)".

Returns

The list of text objects for the numbers.

5.14.5.6 plot_OCV_from_CC_CV() `def ep_bolfi.utility.visualization.plot_OCV_from_CC_CV (`
`ax_ICA_meas,`
`ax_ICA_mean,`
`ax_OCV_meas,`
`ax_OCV_mean,`
`charge,`
`cv,`
`discharge,`
`name,`
`phases,`
`eval_points = 200,`
`spline_SOC_range = (0.01, 0.99),`
`spline_order = 2,`
`spline_smoothing = 2e-3,`
`spline_print = None,`
`parameters_print = False)`

Visualizes the `OCV_fitting.OCV_from_CC_CV` output.

Parameters

<i>ax_ICA_meas</i>	The Axes where the Incremental Capacity Analysis of the measured charge and discharge cycle(s) shall be plotted.
<i>ax_ICA_mean</i>	The Axes where the Incremental Capacity Analysis of the mean voltages of charge and discharge cycle(s) shall be plotted.
<i>ax_OCV_meas</i>	The Axes where the measured voltage curves shall be plotted.
<i>ax_OCV_mean</i>	The Axes where the mean voltage curves shall be plotted.
<i>charge</i>	A <code>Cycling_Information</code> object containing the constant charge cycle(s). If more than one CC-CV-cycle shall be analyzed, please make sure that the order of this, <i>cv</i> and <i>discharge</i> align.
<i>cv</i>	A <code>Cycling_Information</code> object containing the constant voltage part between charge and discharge cycle(s).
<i>discharge</i>	A <code>Cycling_Information</code> object containing the constant discharge cycle(s). These occur after each <i>cv</i> cycle.
<i>name</i>	Name of the material for which the CC-CV-cycling was measured.
<i>phases</i>	Number of phases in the fitting_functions.OCV_fit_function as an int. The higher it is, the more (over-)fitted the model becomes.
<i>eval_points</i>	The number of points for plotting of the OCV curves.
<i>spline_SOC_range</i>	2-tuple giving the SOC range in which the inverted fitting_functions.OCV_fit_function will be interpolated by a smoothing spline. Outside of this range the spline is used for extrapolation. Use this to fit the SOC range of interest more precisely, since a fit of the whole range usually fails due to the singularities at SOC 0 and 1. Please note that this range considers the 0-1-range in which the given SOC lies and not the linear transformation of it from the fitting process.
<i>spline_order</i>	Order of this smoothing spline. If it is set to 0, only the fitting_functions.OCV_fit_function is calculated and plotted.
<i>spline_smoothing</i>	Smoothing factor for this smoothing spline. Default: 2e-3. Lower numbers give more precision, while higher numbers give a simpler spline that smoothes over steep steps in the fitted OCV curve.
<i>spline_print</i>	If set to either 'python' or 'matlab', a string representation of the smoothing spline is printed in the respective format.
<i>parameters_print</i>	Set to True if the fit parameters should be printed to console.

5.14.5.7 push_apart_text() `def ep_bolfi.utility.visualization.push_apart_text (`
`fig,`
`ax,`
`text_objects,`
`lock_xaxis = False,`
`temp_path = "/temp_render.png")`

Push apart overlapping texts until no overlaps remain.

Parameters

<i>fig</i>	The figure which contains the text.
<i>ax</i>	The axis which contains the text.
<i>text_objects</i>	A list of the text objects that shall be pushed apart.
<i>lock_xaxis</i>	If True, texts will only be moved in the y-direction.
<i>temp_path</i>	The path to which a temporary image of the figure "fig" gets saved. This is necessary to establish the text bbox sizes.

5.14.5.8 set_fontsize() `def ep_bolfi.utility.visualization.set_fontsize (`
`ax,`
`title = 12,`
`xaxis = 12,`
`yaxis = 12,`
`xticks = 12,`
`yticks = 12,`
`legend = 12)`

Convenience function for fontsize changes.

Parameters

<i>ax</i>	The axis which texts shall be adjusted.
<i>title</i>	The new fontsize for the title.
<i>xaxis</i>	The new fontsize for the x-axis label.
<i>yaxis</i>	The new fontsize for the y-axis label.
<i>xticks</i>	The new fontsize for the ticks/numbers at the x-axis.
<i>yticks</i>	The new fontsize for the ticks/numbers at the y-axis.
<i>legend</i>	The new fontsize for the legend entries.

5.14.5.9 update_legend() `def ep_bolfi.utility.visualization.update_legend (`
`ax,`
`additional_handles = [],`
`additional_labels = [],`
`additional_handler_map = {})`

Makes sure that all items remain and all items show up.

This basically replaces "ax.legend()" in a way that makes sure that new items can be added to the legend without losing old ones. Please note that only "handler_map"s with class keys work correctly.

Parameters

<i>ax</i>	The axis which legend shall be updated.
<i>additional_handles</i>	The same input "ax.legend(...)" would expect. A list of artists.
<i>additional_labels</i>	The same input "ax.legend(...)" would expect. A list of strings.
<i>additonal_handler_map</i>	The same input "ax.legend(...)" would expect for "handler_map". Please note that, due to the internal structure of the Legend class, only entries with keys that represent classes work right. Entries that have <i>instances</i> of classes (i.e., objects) for keys work exactly once, since the original handle of them is lost in the initialization of a Legend.

5.14.5.10 update_limits() `def ep_bolfi.utility.visualization.update_limits (`

```

    ax,
    xmin = float('inf'),
    xmax = -float('inf'),
    ymin = float('inf'),
    ymax = -float('inf'))

```

Convenience function for adjusting the view.

Parameters

<i>ax</i>	The axis which viewport shall be adjusted.
<i>xmin</i>	The highest lower bound for the x-axis.
<i>xmax</i>	The lowest upper bound for the x-axis.
<i>ymin</i>	The highest lower bound for the y-axis.
<i>ymax</i>	The lowest upper bound for the y-axis.

5.14.5.11 visualize_correlation() `def ep_bolfi.utility.visualization.visualize_correlation (`

```

    fig,
    ax,
    correlation,
    names = None,
    title = None,
    cmap = plt.get_cmap('BrBG'),
    entry_color = 'w')

```

Produces a heatmap of a correlation matrix.

Parameters

<i>fig</i>	The matplotlib.Figure object for plotting.
<i>ax</i>	The matplotlib.Axes object for plotting.
<i>correlation</i>	A two-dimensional (numpy) array that is the correlation matrix.
<i>names</i>	A list of strings that are names of the variables corresponding to each row or column in the correlation matrix.

Parameters

<i>title</i>	The title of the heatmap.
<i>cmap</i>	The matplotlib colormap for the heatmap.
<i>entry_color</i>	The colour of the correlation matrix entries.

6 Class Documentation

6.1 `ep_bolfi.utility.dataset_formatting.Cycling_Information` Class Reference

Contains basic cycling informations.

Inheritance diagram for `ep_bolfi.utility.dataset_formatting.Cycling_Information`:

Collaboration diagram for `ep_bolfi.utility.dataset_formatting.Cycling_Information`:

Public Member Functions

- `def __init__` (self, [timepoints](#), [currents](#), [voltages](#), [other_columns](#)={}, [indices](#)=None)
- `def __len__` (self)
- `def to_json` (self)
- `def from_json` (cls, json_string)
- `def table_descriptors` (self)
- `def table_mapping` (cls)
- `def segment_tables` (self, start=0, stop=None, step=1)
- `def example_table_row` (self)
- `def subslice` (self, start, stop, step=1)
Selects a subslice of the data segments.
- `def subarray` (self, array)
Selects the data segments with the given indices.
- `def extend` (self, other)

Public Attributes

- [timepoints](#)
The times at which measurements were taken.
- [currents](#)
The measured current at those times.
- [voltages](#)
The measured voltage at those times.
- [other_columns](#)
The contents of any other columns.
- [indices](#)
The indices of the individual segments.

6.1.1 Detailed Description

Contains basic cycling informations.

Each member variable is a list and has the same length as the other ones.

6.1.2 Constructor & Destructor Documentation

6.1.2.1 `__init__()` `def ep_bolfi.utility.dataset_formatting.Cycling_Information.__init__(
self,
timepoints,
currents,
voltages,
other_columns = {},
indices = None)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Static_Information](#).

6.1.3 Member Function Documentation

6.1.3.1 `__len__()` `def ep_bolfi.utility.dataset_formatting.Cycling_Information.__len__(
self)`

6.1.3.2 `example_table_row()` `def ep_bolfi.utility.dataset_formatting.Cycling_Information.example_table_row(
self)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.1.3.3 `extend()` `def ep_bolfi.utility.dataset_formatting.Cycling_Information.extend(
self,
other)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

Reimplemented in [ep_bolfi.utility.dataset_formatting.Static_Information](#).

6.1.3.4 from_json() def ep_bolfi.utility.dataset_formatting.Cycling_Information.from_json (
 cls,
 json_string)

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.1.3.5 segment_tables() def ep_bolfi.utility.dataset_formatting.Cycling_Information.segment_tables (
 self,
 start = 0,
 stop = None,
 step = 1)

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.1.3.6 subarray() def ep_bolfi.utility.dataset_formatting.Cycling_Information.subarray (
 self,
 array)

Selects the data segments with the given indices.

Parameters

<i>array</i>	The indices of the segments that are to be returned.
--------------	--

Returns

A new [Cycling_Information](#) object containing the subset.

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

Reimplemented in [ep_bolfi.utility.dataset_formatting.Static_Information](#).

6.1.3.7 subslice() def ep_bolfi.utility.dataset_formatting.Cycling_Information.subslice (
 self,
 start,
 stop,
 step = 1)

Selects a subslice of the data segments.

The arguments exactly match the slice(...) notation.

Parameters

<i>start</i>	The index of the first segment to be included.
<i>stop</i>	The index of the first segment to not be included.
<i>step</i>	Steps between selected segments. Default: 1.

Returns

A new [Cycling_Information](#) object containing the slice.

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

Reimplemented in [ep_bolfi.utility.dataset_formatting.Static_Information](#).

6.1.3.8 table_descriptors() `def ep_bolfi.utility.dataset_formatting.Cycling_Information.table_descriptors (self)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.1.3.9 table_mapping() `def ep_bolfi.utility.dataset_formatting.Cycling_Information.table_mapping (cls)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.1.3.10 to_json() `def ep_bolfi.utility.dataset_formatting.Cycling_Information.to_json (self)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

Reimplemented in [ep_bolfi.utility.dataset_formatting.Static_Information](#).

6.1.4 Member Data Documentation

6.1.4.1 currents `ep_bolfi.utility.dataset_formatting.Cycling_Information.currents`

The measured current at those times.

A list which usually contains lists for segments with variable current and floats for segments with constant current.

6.1.4.2 indices `ep_bolfi.utility.dataset_formatting.Cycling_Information.indices`

The indices of the individual segments.

Defaults to a simple numbering of the segments present. May be used for plotting purposes, e.g., for colourcoding the segments by cycle.

6.1.4.3 other_columns ep_bolfi.utility.dataset_formatting.Cycling_Information.other_columns

The contents of any other columns.

A dictionary ("columns") which values are lists which contain lists for segments. The keys should match user input for the columns.

6.1.4.4 timepoints ep_bolfi.utility.dataset_formatting.Cycling_Information.timepoints

The times at which measurements were taken.

Usually a list of lists where each list corresponds to a segment of the measurement.

6.1.4.5 voltages ep_bolfi.utility.dataset_formatting.Cycling_Information.voltages

The measured voltage at those times.

A list which usually contains lists for segments with variable voltage and floats for segments with constant voltage.

The documentation for this class was generated from the following file:

- [utility/dataset_formatting.py](#)

6.2 ep_bolfi.models.assess_effective_parameters.Effective_Parameters Class Reference

Calculates, stores and prints effective parameters.

Public Member Functions

- `def __init__ (self, parameters, working_electrode='both')`
Preprocesses the model parameters.
- `def eval (self, expression)`
Short-hand for PyBaMM symbol evaluation.
- `def print (self, c_rates=[0.1, 0.2, 0.5, 1.0])`
Prints the voltage losses for the given C-rates.

Public Attributes

- [parameters](#)
The parameter dictionary converted to PyBaMM form.
- [working_electrode](#)
Sets whether full- or half-cell parameters are calculated.
- Q_n
Negative electrode theoretical capacity.
- Q_p
Positive electrode theoretical capacity.
- $R_{c_e_0_1}$
Integration constant for the electrolyte concentration / I.
- $R_{bar_c_n_1}$
Electrolyte concentration / I at the negative electrode.
- $R_{bar_c_{ep}_1}$
Electrolyte concentration / I at the positive electrode.
- $R_{bar_{\phi_{sn}_1}$
Effective resistance of the negative electrode.
- $R_{bar_{\phi_{sp}_1}$
Effective resistance of the positive electrode.
- R_{ns}
Negative electrode resistance (as calculated by the SPMe(S)).
- R_{ps}
Positive electrode resistance (as calculated by the SPMe(S)).
- R_e
Electrolyte resistance (as calculated by the SPMe(S)).
- i_{sn}
SPM(e) negative electrode exchange-current.
- i_{sep}
SPM(e) positive electrode exchange-current.

6.2.1 Detailed Description

Calculates, stores and prints effective parameters.

6.2.2 Constructor & Destructor Documentation

6.2.2.1 `__init__()` `def ep_bolfi.models.assess_effective_parameters.Effective_Parameters.__init__(
self,
parameters,
working_electrode = 'both')`

Preprocesses the model parameters.

Parameters

<i>parameters</i>	A dictionary of parameter values with the parameter names as keys. For these names, please refer to ep_bolfi.models.standard_parameters .
<i>working_electrode</i>	When set to either 'negative' or 'positive', the parameters will be treated as a half-cell setup with said electrode.

6.2.3 Member Function Documentation

6.2.3.1 eval() def ep_bolfi.models.assess_effective_parameters.Effective_Parameters.eval (
 self,
 expression)

Short-hand for PyBaMM symbol evaluation.

Parameters

<i>expression</i>	A pybamm.Symbol.
-------------------	------------------

Returns

The numeric value of "expression".

6.2.3.2 print() def ep_bolfi.models.assess_effective_parameters.Effective_Parameters.print (
 self,
 c_rates = [0.1, 0.2, 0.5, 1.0])

Prints the voltage losses for the given C-rates.

Parameters

<i>c_rates</i>	The C-rates (as fraction of "Typical current [A]").
----------------	---

6.2.4 Member Data Documentation

6.2.4.1 i_{se} ep_bolfi.models.assess_effective_parameters.Effective_Parameters. i_{se}

SPM(e) negative electrode exchange-current.

6.2.4.2 i_{sep} ep_bolfi.models.assess_effective_parameters.Effective_Parameters. i_{sep}

SPM(e) positive electrode exchange-current.

6.2.4.3 **parameters** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.parameters`

The parameter dictionary converted to PyBaMM form.

Convert SubstitutionDict to dict by iterating over it.

6.2.4.4 **Q_n** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.Q_n`

Negative electrode theoretical capacity.

6.2.4.5 **Q_p** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.Q_p`

Positive electrode theoretical capacity.

6.2.4.6 **$R_{bar_c_n_1}$** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.R_bar_c_n_1`

Electrolyte concentration / I at the negative electrode.

6.2.4.7 **$R_{bar_c_p_1}$** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.R_bar_c_p_1`

Electrolyte concentration / I at the positive electrode.

6.2.4.8 **$R_{bar_phi_n_1}$** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.R_bar_phi_n_1`

Effective resistance of the negative electrode.

6.2.4.9 **$R_{bar_phi_p_1}$** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.R_bar_phi_p_1`

Effective resistance of the positive electrode.

6.2.4.10 **$R_{c_e_0_1}$** `ep_bolfi.models.assess_effective_parameters.Effective_Parameters.R_c_e_0_1`

Integration constant for the electrolyte concentration / I.

6.2.4.11 R_e ep_bolfi.models.assess_effective_parameters.Effective_Parameters. R_e

Electrolyte resistance (as calculated by the SPMe(S)).

6.2.4.12 R_{ns} ep_bolfi.models.assess_effective_parameters.Effective_Parameters. R_{ns}

Negative electrode resistance (as calculated by the SPMe(S)).

6.2.4.13 R_{ps} ep_bolfi.models.assess_effective_parameters.Effective_Parameters. R_{ps}

Positive electrode resistance (as calculated by the SPMe(S)).

6.2.4.14 `working_electrode` ep_bolfi.models.assess_effective_parameters.Effective_Parameters.working_electrode

Sets whether full- or half-cell parameters are calculated.

The documentation for this class was generated from the following file:

- [models/assess_effective_parameters.py](#)

6.3 ep_bolfi.models.electrolyte.Electrolyte Class Reference

[Electrolyte](#) model assuming a symmetric Li-metal cell.

Inheritance diagram for ep_bolfi.models.electrolyte.Electrolyte:

6.4 ep_bolfi.models.electrolyte.Electrolyte_internal Class Reference

Defining equations for a symmetric Li cell with electrolyte.

Inheritance diagram for ep_bolfi.models.electrolyte.Electrolyte_internal:

Collaboration diagram for ep_bolfi.models.electrolyte.Electrolyte_internal:

Public Member Functions

- `def __init__ (self, param, pybamm_control=False, options={}, build=True)`
Sets the model properties.
- `def get_fundamental_variables (self)`
Builds all relevant model variables' symbols.
- `def get_coupled_variables (self, variables)`
Builds all model symbols that rely on other models.
- `def set_rhs (self, variables)`
Sets up the right-hand-side equations in `self.rhs`.
- `def set_algebraic (self, variables)`
Sets up the algebraic equations in `self.algebraic`.
- `def set_boundary_conditions (self, variables)`
Sets the (self.)`boundary`(_)conditions.
- `def set_initial_conditions (self, variables)`
Sets the (self.)`initial`(_)conditions.
- `def set_events (self, variables)`
Sets up the termination switches in `self.events`.

Public Attributes

- [pybamm_control](#)
Current is fixed if False and a variable if True.

6.4.1 Detailed Description

Defining equations for a symmetric Li cell with electrolyte.

6.4.1.1 Reference SG Marquis, V Sulzer, R Timms, CP Please and SJ Chapman. “An asymptotic derivation of a single particle model with electrolyte”. Journal of The Electrochemical Society, 166(15):A3693–A3706, 2019

6.4.2 Constructor & Destructor Documentation

6.4.2.1 `__init__()` `def ep_bolfi.models.electrolyte.Electrolyte_internal.__init__ (`
`self,`
`param,`
`pybamm_control = False,`
`options = {},`
`build = True)`

Sets the model properties.

Parameters

<i>param</i>	A class containing all the relevant parameters for this model. For example, models.standard_parameters represents a valid choice for this parameter.
<i>pybamm_control</i>	Per default False, which indicates that the current is given as a function. If set to True, this model is compatible with PyBaMM experiments, e.g. CC-CV simulations. The current is then a variable and it or voltage can be fixed functions. Generated by Doxygen
<i>options</i>	Not used; only here for compatibility with the base class.
<i>build</i>	Not used; only here for compatibility with the base class.

6.4.3 Member Function Documentation

6.4.3.1 get_coupled_variables() `def ep_bolfi.models.electrolyte.Electrolyte_internal.get_coupled_variables (self, variables)`

Builds all model symbols that rely on other models.

Parameters

<i>variables</i>	A dictionary containing at least all variable symbols that are required for the variable symbols built here.
------------------	--

Returns

A dictionary with the new variables' names as keys and their symbols (of type pybamm.Symbol) as values.

6.4.3.2 get_fundamental_variables() `def ep_bolfi.models.electrolyte.Electrolyte_internal.get_fundamental_variables (self)`

Builds all relevant model variables' symbols.

Returns

A dictionary with the variables' names as keys and their symbols (of type pybamm.Symbol) as values.

6.4.3.3 set_algebraic() `def ep_bolfi.models.electrolyte.Electrolyte_internal.set_algebraic (self, variables)`

Sets up the algebraic equations in self.algebraic.

6.4.3.4 set_boundary_conditions() `def ep_bolfi.models.electrolyte.Electrolyte_internal.set_boundary_conditions (self, variables)`

Sets the (self.)boundary(_)conditions.

6.4.3.5 set_events() def ep_bolfi.models.electrolyte.Electrolyte_internal.set_events (
 self,
 variables)

Sets up the termination switches in self.events.

6.4.3.6 set_initial_conditions() def ep_bolfi.models.electrolyte.Electrolyte_internal.set_initial_conditions (
 self,
 variables)

Sets the (self.)initial(_)conditions.

6.4.3.7 set_rhs() def ep_bolfi.models.electrolyte.Electrolyte_internal.set_rhs (
 self,
 variables)

Sets up the right-hand-side equations in self.rhs.

6.4.4 Member Data Documentation

6.4.4.1 pybamm_control ep_bolfi.models.electrolyte.Electrolyte_internal.pybamm_control

Current is fixed if False and a variable if True.

The documentation for this class was generated from the following file:

- models/[electrolyte.py](#)

6.5 ep_bolfi.optimization.EP_BOLFI.EP_BOLFI Class Reference

Expectation Propagation and Bayesian Optimization.

Public Member Functions

- def [__init__](#) (self, [simulators](#), [experimental_datasets](#), [feature_extractors](#), [fixed_parameters](#), [free_parameters](#)=None, [initial_covariance](#)=None, [free_parameters_boundaries](#)=None, [boundaries_in_deviations](#)=0, [Q](#)=None, [r](#)=None, [Q_features](#)=None, [r_features](#)=None, [transform_parameters](#)={}, [weights](#)=None, [display_current_feature](#)=None, [fixed_parameter_order](#)=None)
- def [result_to_json](#) (self, seed=None)
- def [log_to_json](#) (self)
- def [visualize_parameter_distribution](#) (self)
Plots the features and visualizes the correlation.
- def [run](#) (self, [bolfi_initial_evidence](#)=None, [bolfi_total_evidence](#)=None, [bolfi_posterior_samples](#)=None, [ep_iterations](#)=3, [ep_dampener](#)=None, [final_dampening](#)=None, [ep_dampener_reduction_steps](#)=-1, [gelman_rubin_threshold](#)=None, [ess_ratio_resample](#)=5.0, [ess_ratio_sampling_from_zero](#)=-1.0, [ess_ratio_abort](#)=20.0, [max_heuristic_steps](#)=10, [posterior_sampling_increase](#)=1.2, [model_resampling_increase](#)=1.0, [independent_mcmc_chains](#)=4, [scramble_ep_feature_order](#)=True, [show_trials](#)=False, [verbose](#)=True, seed=None)
Runs Expectation Propagation together with BOLFI.

Public Attributes

- [simulators](#)
- [experimental_datasets](#)
- [feature_extractors](#)
- [fixed_parameters](#)
- [free_parameters](#)
- [initial_covariance](#)
- [free_parameters_boundaries](#)
- [boundaries_in_deviations](#)
Uses boundaries as multiples of the standard deviation.
- [transform_parameters](#)
- [weights](#)
Set the weights to unity if None are given.
- [display_current_feature](#)
- [fixed_parameter_order](#)
If the parameter order is explicitly given, use that instead.
- [log_of_tried_parameters](#)
Stores all parameter combinations that have been tried.
- [experimental_features](#)
Experimental features.
- [input_dim](#)
Input dimension of the estimation task.
- [output_dim](#)
Output dimension of the estimation task (sum of features).
- [simulator_index_by_feature](#)
Mapping of index by all features to corresponding simulator.
- [sub_index_by_feature](#)
Mapping of index by all features to that by one set of them.
- [initial_guesses](#)
Container for the initial expectation values.
- [log_of_raw_tried_parameters](#)
Stores all raw parameter evaluation points.
- [log_of_discrepancies](#)
Stores all discrepancies of the sampled parameters.
- [final_expectation](#)
Stores the inference mean (empty at first).
- [final_covariance](#)
Stores the inference covariance (empty at first).
- [final_correlation](#)
Stores the inference correlation (empty at first).
- [initial_Q](#)
Expectation Propagation covariance matrix (prior).
- [Q](#)
Expectation Propagation covariance matrix (posterior).
- [initial_r](#)
Expectation Propagation expectation value (prior).
- [r](#)
Expectation Propagation expectation value (posterior).
- [Q_features](#)
Expectation Propagation itemized covariance matrices.
- [r_features](#)

Expectation Propagation itemized expectation values.

- [inferred_parameters](#)

The inferred model parameters.

- [final_error_bounds](#)

The 95% confidence bounds (which don't reflect the cross-correlations, but are easier to interpret).

6.5.1 Detailed Description

Expectation Propagation and Bayesian Optimization.

Sets up and runs these two algorithms to infer model parameters. Use the variables "Q", "r", "Q_features" and "r_features" to copy the state of another estimator. Do not use them in any other case. Always use either all of them or none of them.

6.5.2 Constructor & Destructor Documentation

6.5.2.1 `__init__()` `def ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.__init__(`
`self,`
`simulators,`
`experimental_datasets,`
`feature_extractors,`
`fixed_parameters,`
`free_parameters = None,`
`initial_covariance = None,`
`free_parameters_boundaries = None,`
`boundaries_in_deviations = 0,`
`Q = None,`
`r = None,`
`Q_features = None,`
`r_features = None,`
`transform_parameters = {},`
`weights = None,`
`display_current_feature = None,`
`fixed_parameter_order = None)`

Parameters

<i>simulators</i>	A list of functions that take one argument: a dictionary of combined 'fixed_parameters' and 'free_parameters'. They return the simulated counterpart to the experimental data. Most of the time, one function will be sufficient. Additional functions may be used to combine simulators which each give a subset of the total experimental method.
<i>experimental_datasets</i>	A list of the experimental data. Each entry corresponds to the simulator in 'simulators' with the same index and has the same structure as its output.
<i>feature_extractors</i>	A list of functions which each take the corresponding data entry and return a list of numbers, which represent its reduced features.
<i>fixed_parameters</i>	Dictionary of parameters that stay fixed and their values.

Parameters

<i>free_parameters</i>	Dictionary of parameters which shall be inferred and their initial guesses or, more accurately, their expected values. Please note that these values live in the transformed space. Optionally, the values may be a 2-tuple where the second entry would be the variance of that parameter. For finer tuning with covariances, use 'initial_covariance' (will take precedence). Alternatively, you may set 'free_parameters_boundaries' to set the expected values and variances by confidence intervals.
<i>initial_covariance</i>	Initial covariance of the parameters. Has to be a symmetric matrix (list of list or numpy 2D array). A reasonable simple choice is to have a diagonal matrix and set the standard deviation σ_i of each parameter to half of the distance between initial guess and biggest/smallest value that shall be tried. If the diagonal entries are σ_i^2 and the bounds are symmetric, the probability distribution of each parameter is 95% within these bounds. Please note that the same does not hold for the whole probability distribution.
<i>free_parameters_boundaries</i>	Optional hard boundaries of the space in which optimal parameters are searched for. They are given as a dictionary with values as 2-tuples with the left and right boundaries. Boundaries need to be given for either none or all parameters. If None are given, boundaries will be set by 'boundaries_in_deviations' relative to the covariance. The default then is the 95 % confidence ellipsoid. If neither 'initial_covariance' nor 'free_parameters' set the covariance, this parameter sets it according to the example in the description of 'initial_covariance'.
<i>boundaries_in_deviations</i>	When ≤ 0 , the boundaries are set as described above. When > 0 , the boundaries are this multiple of the standard deviation. This scales with the shrinking covariance as the algorithm progresses. 'free_parameters_boundaries' takes precedence, i.e., the covariance gets set and then the boundaries in the optimization are in standard deviations.
<i>Q</i>	If you want to restore a previous EP-BOLFI instance from a dump of its data (see "result_to_json" method), put the "Q" attribute stored therein into this parameter.
<i>r</i>	Same as Q, but use the "r" attribute.
<i>Q_features</i>	Same as Q, but use the "Q_features" attribute.
<i>r_features</i>	Same as Q, but use the "r_features" attribute.
<i>transform_parameters</i>	Optional transformations between the parameter space that is used for searching for optimal parameters and the model parameters. Any missing free parameter is not transformed. The values are 2-tuples. The first entry is a function taking the search space parameter and returning the model parameter. The second entry is the inverse function. For convenience, any value may also be one of the following: <ul style="list-style-type: none"> 'none' => (identity, identity) 'log' => (exp, log) Please note that, for performance reasons, the returned inferred values are directly back-transformed from the mean of the internal standard distribution. This means that they represent the median of the actual distribution.
<i>weights</i>	Optional weights to assign to the features. Higher weights give more importance to a feature and vice versa. A list of lists which correspond to the 'feature_extractors'.
<i>display_current_feature</i>	A list of functions. Each corresponds to a feature extractor with the same index. Given an index of the array of its features, this returns a short description of it. If None is given, only the index will be shown in the output.

Parameters

<i>fixed_parameter_order</i>	Establish a numerical order to the parameter names. This prevents errors arising from internal reordering of the dictionaries. Only necessary when using the same model in different contexts.
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6.5.3 Member Function Documentation

6.5.3.1 log_to_json() `def ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.log_to_json (self)`

6.5.3.2 result_to_json() `def ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.result_to_json (self, seed = None)`

6.5.3.3 run() `def ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.run (self, bolfi_initial_evidence = None, bolfi_total_evidence = None, bolfi_posterior_samples = None, ep_iterations = 3, ep_dampener = None, final_dampening = None, ep_dampener_reduction_steps = -1, gelman_rubin_threshold = None, ess_ratio_resample = 5.0, ess_ratio_sampling_from_zero = -1.0, ess_ratio_abort = 20.0, max_heuristic_steps = 10, posterior_sampling_increase = 1.2, model_resampling_increase = 1.1, independent_mcmc_chains = 4, scramble_ep_feature_order = True, show_trials = False, verbose = True, seed = None)`

Runs Expectation Propagation together with BOLFI.

This function can be called multiple times; the estimation will take off from where it last stopped.

Parameters

<i>bolfi_initial_evidence</i>	Number of evidence samples BOLFI will take for each feature before using Bayesian Optimization sampling. Default: $1 + 2^{\text{number of estimated parameters}}$.
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Parameters

<i>bolfi_total_evidence</i>	Number of evidence samples BOLFI will take for each feature in total (including initial evidence). Default: $2 * \text{"bolfi_initial_evidence"}$.
<i>bolfi_posterior_samples</i>	Effective number of samples BOLFI will take from the posterior distribution. These are then used to fit a Gaussian to the posterior. Fit convergence scales with $1/\sqrt{n}$. Default: $I^2 + 3 * I$ with I as the number of estimated parameters. This is the number of the metaparameters of the underlying probability distribution times 2. The "times 2" considers the warmup samples.
<i>ep_iterations</i>	The number of iterations of the Expectation Propagation algorithm, i.e., the number of passes over each feature. Default: 3.
<i>ep_dampener</i>	The linear combination factor of the posterior calculated by BOLFI and the pseudo-prior. 0 means no dampening, i.e., the pseudo-prior gets replaced by the posterior. For values up to 1, that fraction of the pseudo-prior remains in each site update. Default: with "a" as the number of features and "b" as "ep_iterations", $1 - a * (1 - \sqrt[b]{\text{"final_dampening"}})$.
<i>final_dampening</i>	Alternative way to set "ep_dampener". 0 means no dampening. For values up to 1, that fraction of the prior remains after the whole estimation. Default: if "ep_dampener" is not set, 0.5. Else, "ep_dampener" takes precedence.
<i>ep_dampener_reduction_steps</i>	Number of iterations over which the 'ep_dampener' gets reduced to 0. In each iteration, an equal fraction of it gets subtracted. Set to a negative number to disable the reduction. Default: -1.
<i>gelman_rubin_threshold</i>	Optional threshold on top of the effective sample size. Values close to one indicate a converged estimate of the pseudo-posteriors. Never set to exactly one.
<i>ess_ratio_sampling_from_zero</i>	Threshold in the ratio of effective sample size to samples in the pseudo-posterior estimation, at which the sampling defaults to starting at the center of the pseudo-prior. Set higher than "ess_ratio_resample" to disable this behaviour.
<i>ess_ratio_resample</i>	Threshold in the ratio of effective sample size to samples in the pseudo-posterior estimation, at which before sampling the model gets resampled. Set higher than "ess_ratio_abort" to disable this behaviour.
<i>ess_ratio_abort</i>	Threshold in the ratio of effective sample size to samples in the pseudo-posterior estimation, at which the sampling aborts and the pseudo-posterior update is skipped.
<i>max_heuristic_steps</i>	The heuristics that are set by the "ess_ratio_*" arguments could effectively run forever. This parameter limits the amount of times these heuristics get employed in on EP iteration before it terminates.
<i>posterior_sampling_increase</i>	The factor by which the ratio of the effective sample size to samples in the pseudo-posterior estimation is multiplied each loop (cumulatively). Never set to exactly one or lower, as it might result in an infinite loop.
<i>model_resampling_increase</i>	The factor by which 'bolfi_total_evidence' gets multiplied each time the model gets resampled.
<i>independent_mcmc_chains</i>	The number of independent Markov-Chain Monte Carlo chains that are used for the estimation of the pseudo-posterior. Since we did not implement parallelization, more chains will not be faster, but more stable.
<i>scramble_ep_feature_order</i>	True randomizes the order that the EP features are iterated over. Their order is still consistent across EP iterations. False uses the order that the "feature_extractors" define.
<i>show_trials</i>	True plots the log of tried parameters live. Please note that each plot blocks the execution of the program, so do not use this when running the estimation in the background.

Parameters

<i>verbose</i>	True shows verbose error messages and logs of the estimation process. With False, you need to get the estimation results from <code>self.final_expectation</code> and <code>self.final_covariance</code> . Default: True.
<i>seed</i>	Optional seed that is used in the RNG. If None is given, the results will be slightly different each time.

Returns

The BOLFI instance of the last EP iteration. As such, it contains the Posterior of the overall inference procedure.

6.5.3.4 visualize_parameter_distribution() `def ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.visualize_parameter_↔
distribution (
self)`

Plots the features and visualizes the correlation.

Please note that this function requires that the output of the individual simulators and the individual experimental data give an x- and y-axis when indexed with [0] and [1], respectively. Lists of lists in [0] and [1] with segmented data works as well. [EP_BOLFI.run\(\)](#) does not have these restrictions. Visualizes the comparison that [EP_BOLFI](#) was set up to infer model parameters with. May be used to check if everything works as intended. Additionally, the 95% confidence error bounds for the parameters are visualized to check for reasonable bounds. If called after '[run\(\)](#)', the expected parameter set, the correlation and error bounds are from the finished estimation.

6.5.4 Member Data Documentation

6.5.4.1 boundaries_in_deviations `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.boundaries_in_deviations`

Uses boundaries as multiples of the standard deviation.

6.5.4.2 display_current_feature `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.display_current_feature`

6.5.4.3 experimental_datasets `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.experimental_datasets`

6.5.4.4 experimental_features `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.experimental_features`

Experimental features.

6.5.4.5 `feature_extractors` `ep_bolfi.optimization.EP_BOLFI.feature_extractors`

6.5.4.6 `final_correlation` `ep_bolfi.optimization.EP_BOLFI.final_correlation`

Stores the inference correlation (empty at first).

6.5.4.7 `final_covariance` `ep_bolfi.optimization.EP_BOLFI.final_covariance`

Stores the inference covariance (empty at first).

6.5.4.8 `final_error_bounds` `ep_bolfi.optimization.EP_BOLFI.final_error_bounds`

The 95% confidence bounds (which don't reflect the cross-correlations, but are easier to interpret).

6.5.4.9 `final_expectation` `ep_bolfi.optimization.EP_BOLFI.final_expectation`

Stores the inference mean (empty at first).

6.5.4.10 `fixed_parameter_order` `ep_bolfi.optimization.EP_BOLFI.fixed_parameter_order`

If the parameter order is explicitly given, use that instead.

6.5.4.11 `fixed_parameters` `ep_bolfi.optimization.EP_BOLFI.fixed_parameters`

6.5.4.12 `free_parameters` `ep_bolfi.optimization.EP_BOLFI.free_parameters`

6.5.4.13 `free_parameters_boundaries` `ep_bolfi.optimization.EP_BOLFI.free_parameters_boundaries`

6.5.4.14 inferred_parameters `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.inferred_parameters`

The inferred model parameters.

6.5.4.15 initial_covariance `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.initial_covariance`**6.5.4.16 initial_guesses** `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.initial_guesses`

Container for the initial expectation values.

6.5.4.17 initial_Q `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.initial_Q`

Expectation Propagation covariance matrix (prior).

6.5.4.18 initial_r `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.initial_r`

Expectation Propagation expectation value (prior).

6.5.4.19 input_dim `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.input_dim`

Input dimension of the estimation task.

6.5.4.20 log_of_discrepancies `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.log_of_discrepancies`

Stores all discrepancies of the sampled parameters.

6.5.4.21 log_of_raw_tried_parameters `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.log_of_raw_tried_parameters`

Stores all raw parameter evaluation points.

6.5.4.22 log_of_tried_parameters `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.log_of_tried_parameters`

Stores all parameter combinations that have been tried.

6.5.4.23 `output_dim` `ep_bolfi.optimization.EP_BOLFI.output_dim`

Output dimension of the estimation task (sum of features).

6.5.4.24 `Q` `ep_bolfi.optimization.EP_BOLFI.Q`

Expectation Propagation covariance matrix (posterior).

6.5.4.25 `Q_features` `ep_bolfi.optimization.EP_BOLFI.Q_features`

Expectation Propagation itemized covariance matrices.

6.5.4.26 `r` `ep_bolfi.optimization.EP_BOLFI.r`

Expectation Propagation expectation value (posterior).

6.5.4.27 `r_features` `ep_bolfi.optimization.EP_BOLFI.r_features`

Expectation Propagation itemized expectation values.

6.5.4.28 `simulator_index_by_feature` `ep_bolfi.optimization.EP_BOLFI.simulator_index_by_feature`

Mapping of index by all features to corresponding simulator.

6.5.4.29 `simulators` `ep_bolfi.optimization.EP_BOLFI.simulators`

6.5.4.30 `sub_index_by_feature` `ep_bolfi.optimization.EP_BOLFI.sub_index_by_feature`

Mapping of index by all features to that by one set of them.

6.5.4.31 `transform_parameters` `ep_bolfi.optimization.EP_BOLFI.transform_parameters`

6.5.4.32 **weights** `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI.weights`

Set the weights to unity if None are given.

The documentation for this class was generated from the following file:

- optimization/[EP_BOLFI.py](#)

6.6 `ep_bolfi.utility.dataset_formatting.Impedance_Measurement` Class Reference

Contains basic impedance data.

Inheritance diagram for `ep_bolfi.utility.dataset_formatting.Impedance_Measurement`:

Collaboration diagram for `ep_bolfi.utility.dataset_formatting.Impedance_Measurement`:

Public Member Functions

- `def __init__ (self, frequencies, real_impedances, imaginary_impedances, phases, other_columns={}, indices=None)`
- `def __len__ (self)`
- `def complex_impedances (self)`
- `def to_json (self)`
- `def from_json (cls, json_string)`
- `def table_descriptors (self)`
- `def table_mapping (cls)`
- `def segment_tables (self, start=0, stop=None, step=1)`
- `def example_table_row (self)`
- `def subslice (self, start, stop, step=1)`
Selects a subslice of the data segments.
- `def subarray (self, array)`
Selects the data segments with the given indices.
- `def extend (self, other)`

Public Attributes

- `frequencies`
The frequencies at which impedances were measured.
- `real_impedances`
The real part of the impedances measured at those frequencies.
- `imaginary_impedances`
The imaginary part of the impedances measured at those frequencies.
- `phases`
The phases of the impedance measured at those frequencies.
- `other_columns`
The contents of any other columns.
- `indices`
The indices of the individual segments.

6.6.1 Detailed Description

Contains basic impedance data.

Each member variable is a list and has the same length as the other ones.

6.6.2 Constructor & Destructor Documentation

6.6.2.1 `__init__()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.__init__(
 self,
 frequencies,
 real_impedances,
 imaginary_impedances,
 phases,
 other_columns = {},
 indices = None)`

6.6.3 Member Function Documentation

6.6.3.1 `__len__()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.__len__(
 self)`

6.6.3.2 `complex_impedances()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.complex_impedances(
 self)`

6.6.3.3 `example_table_row()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.example_table_row(
 self)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.6.3.4 `extend()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.extend(
 self,
 other)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.6.3.5 from_json() `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.from_json (`
 cls,
 json_string)

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.6.3.6 segment_tables() `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.segment_tables (`
 self,
 start = 0,
 stop = None,
 step = 1)

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.6.3.7 subarray() `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.subarray (`
 self,
 array)

Selects the data segments with the given indices.

Parameters

<i>array</i>	The indices of the segments that are to be returned.
--------------	--

Returns

A new [Impedance_Measurement](#) object containing the subset.

Reimplemented from [ep_bolfi.utility.dataset_formatting.Measurement](#).

6.6.3.8 subslice() `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.subslice (`
 self,
 start,
 stop,
 step = 1)

Selects a subslice of the data segments.

The arguments exactly match the slice(...) notation.

Parameters

<i>start</i>	The index of the first segment to be included.
<i>stop</i>	The index of the first segment to not be included.
<i>step</i>	Steps between selected segments. Default: 1.

Returns

A new `Impedance_Measurement` object containing the slice.

Reimplemented from `ep_bolfi.utility.dataset_formatting.Measurement`.

6.6.3.9 `table_descriptors()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.table_descriptors (self)`

Reimplemented from `ep_bolfi.utility.dataset_formatting.Measurement`.

6.6.3.10 `table_mapping()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.table_mapping (cls)`

Reimplemented from `ep_bolfi.utility.dataset_formatting.Measurement`.

6.6.3.11 `to_json()` `def ep_bolfi.utility.dataset_formatting.Impedance_Measurement.to_json (self)`

Reimplemented from `ep_bolfi.utility.dataset_formatting.Measurement`.

6.6.4 Member Data Documentation

6.6.4.1 `frequencies` `ep_bolfi.utility.dataset_formatting.Impedance_Measurement.frequencies`

The frequencies at which impedances were measured.

Usually a list of lists where each list corresponds to a different equilibrium.

6.6.4.2 `imaginary_impedances` `ep_bolfi.utility.dataset_formatting.Impedance_Measurement.imaginary_impedances`

The imaginary part of the impedances measured at those frequencies.

6.6.4.3 `indices` `ep_bolfi.utility.dataset_formatting.Impedance_Measurement.indices`

The indices of the individual segments.

Defaults to a simple numbering of the segments present. May be used for plotting purposes, e.g., for colourcoding the segments by cycle.

6.6.4.4 **other_columns** `ep_bolfi.utility.dataset_formatting.Impedance_Measurement.other_columns`

The contents of any other columns.

A dictionary ("columns") which values are lists which contain lists for segments. The keys should match user input for the columns.

6.6.4.5 **phases** `ep_bolfi.utility.dataset_formatting.Impedance_Measurement.phases`

The phases of the impedance measured at those frequencies.

6.6.4.6 **real_impedances** `ep_bolfi.utility.dataset_formatting.Impedance_Measurement.real_impedances`

The real part of the impedances measured at those frequencies.

The documentation for this class was generated from the following file:

- [utility/dataset_formatting.py](#)

6.7 **ep_bolfi.utility.dataset_formatting.Measurement Class Reference**

Defines common methods for measurement objects.

Inheritance diagram for `ep_bolfi.utility.dataset_formatting.Measurement`:

Public Member Functions

- `def to_json (self)`
- `def from_json (cls, json_string)`
- `def table_descriptors (self)`
- `def table_mapping (cls)`
- `def segment_tables (self, start=0, stop=None, step=1)`
- `def example_table_row (self)`
- `def subslice (self, start, stop, step=1)`
- `def subarray (self, array)`
- `def extend (self, other)`

6.7.1 Detailed Description

Defines common methods for measurement objects.

6.7.2 Member Function Documentation

6.7.2.1 `example_table_row()` `def ep_bolfi.utility.dataset_formatting.Measurement.example_table_row (`
`self)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), and [ep_bolfi.utility.dataset_formatting.Cycling_In](#)

6.7.2.2 `extend()` `def ep_bolfi.utility.dataset_formatting.Measurement.extend (`
`self,`
`other)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), [ep_bolfi.utility.dataset_formatting.Static_Informa](#)
and [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.7.2.3 `from_json()` `def ep_bolfi.utility.dataset_formatting.Measurement.from_json (`
`cls,`
`json_string)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), and [ep_bolfi.utility.dataset_formatting.Cycling_In](#)

6.7.2.4 `segment_tables()` `def ep_bolfi.utility.dataset_formatting.Measurement.segment_tables (`
`self,`
`start = 0,`
`stop = None,`
`step = 1)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), and [ep_bolfi.utility.dataset_formatting.Cycling_In](#)

6.7.2.5 `subarray()` `def ep_bolfi.utility.dataset_formatting.Measurement.subarray (`
`self,`
`array)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), [ep_bolfi.utility.dataset_formatting.Static_Informa](#)
and [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.7.2.6 `subslice()` `def ep_bolfi.utility.dataset_formatting.Measurement.subslice (`
`self,`
`start,`
`stop,`
`step = 1)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), [ep_bolfi.utility.dataset_formatting.Static_Informa](#)
and [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.7.2.7 table_descriptors() `def ep_bolfi.utility.dataset_formatting.Measurement.table_descriptors (self)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), and [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.7.2.8 table_mapping() `def ep_bolfi.utility.dataset_formatting.Measurement.table_mapping (cls)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), and [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.7.2.9 to_json() `def ep_bolfi.utility.dataset_formatting.Measurement.to_json (self)`

Reimplemented in [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#), [ep_bolfi.utility.dataset_formatting.Static_Information](#), and [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

The documentation for this class was generated from the following file:

- [utility/dataset_formatting.py](#)

6.8 ep_bolfi.optimization.EP_BOLFI.NDArrayEncoder Class Reference

Inheritance diagram for `ep_bolfi.optimization.EP_BOLFI.NDArrayEncoder`:

Collaboration diagram for `ep_bolfi.optimization.EP_BOLFI.NDArrayEncoder`:

Public Member Functions

- `def default (self, item)`

6.8.1 Member Function Documentation

6.8.1.1 default() `def ep_bolfi.optimization.EP_BOLFI.NDArrayEncoder.default (self, item)`

The documentation for this class was generated from the following file:

- [optimization/EP_BOLFI.py](#)

6.9 ep_bolfi.utility.fitting_functions.NDArrayEncoder Class Reference

Inheritance diagram for ep_bolfi.utility.fitting_functions.NDArrayEncoder:

Collaboration diagram for ep_bolfi.utility.fitting_functions.NDArrayEncoder:

Public Member Functions

- def [default](#) (self, item)

6.9.1 Member Function Documentation

6.9.1.1 default() def ep_bolfi.utility.fitting_functions.NDArrayEncoder.default (
 self,
 item)

The documentation for this class was generated from the following file:

- utility/[fitting_functions.py](#)

6.10 ep_bolfi.utility.fitting_functions.OCV_fit_result Class Reference

Contains OCV fit parameters and related information.

Inheritance diagram for ep_bolfi.utility.fitting_functions.OCV_fit_result:

Collaboration diagram for ep_bolfi.utility.fitting_functions.OCV_fit_result:

Public Member Functions

- def [__init__](#) (self, fit, SOC, OCV, SOC_offset=1.0, SOC_scale=1.0, [optimize_result](#)=None, [spline_interpolation_knots](#)=None, [spline_interpolation_coefficients](#)=None, [function_string](#)=None)
- def [to_json](#) (self)
- def [SOC_adjusted](#) (self, soc=None)
 Gives the adjusted SOC values.
- def [SOC_other_electrode](#) (self, soc=None)
 Relates the SOC's of the two electrodes to each other.

Public Attributes

- [SOC_range](#)
The SOC range of the data.
- [fit](#)
The fit parameters of the OCV function from Birkel.
- [E_0](#)
The E_0 (plateau voltages) parameters.
- [a](#)
The a (inverse plateau widths) parameters.
- [Δx](#)
The Δx (phase proportion) parameters.
- [SOC](#)
The SOC data points.
- [OCV](#)
The OCV data points.
- [SOC_offset](#)
If another electrode was factored out in the data, this may contain its SOC at SOC 0 of the electrode of interest.
- [SOC_scale](#)
If another electrode was factored out in the data, this may contain the the rate of change of its SOC to that of the electrode of interest.
- [optimize_result](#)
The `scipy.optimize.OptimizeResult` that led to the fit.
- [spline_interpolation_knots](#)
The knots of the interpolating spline fitted to the inverse.
- [spline_interpolation_coefficients](#)
The coefficients of the interpolating spline fitted to the inverse.
- [function_string](#)
The string representation of the interpolating spline fitted to the inverse.

6.10.1 Detailed Description

Contains OCV fit parameters and related information.

6.10.1.1 Reference C. R. Birkel, E. McTurk, M. R. Roberts, P. G. Bruce and D. A. Howey. “A Parametric Open Circuit Voltage Model for Lithium Ion Batteries”. Journal of The Electrochemical Society, 162(12):A2271-A2280, 2015

6.10.2 Constructor & Destructor Documentation

6.10.2.1 `__init__()` `def ep_bolfi.utility.fitting_functions.OCV_fit_result.__init__(`
`self,`
`fit,`
`SOC,`
`OCV,`
`SOC_offset = 1.0,`
`SOC_scale = 1.0,`
`optimize_result = None,`
`spline_interpolation_knots = None,`
`spline_interpolation_coefficients = None,`
`function_string = None)`

6.10.3 Member Function Documentation

6.10.3.1 SOC_adjusted() `def ep_bolfi.utility.fitting_functions.OCV_fit_result.SOC_adjusted (`
 `self,`
 `soc = None)`

Gives the adjusted SOC values.

Parameters

<code>soc</code>	The SOC as assigned in the original data. This usually corresponds to the range available during a measurement.
------------------	---

Returns

The SOC as corrected by the OCV model. These values will try to correspond to the level of lithiation.

6.10.3.2 SOC_other_electrode() `def ep_bolfi.utility.fitting_functions.OCV_fit_result.SOC_other_electrode (`
 `self,`
 `soc = None)`

Relates the SOC of the two electrodes to each other.

If the original data was of a full cell and the other electrode was factored out, this may contain the function that takes the SOC of the electrode of interest and gives the SOC of the other electrode, i.e., the stoichiometric relation.

Parameters

<code>soc</code>	The SOC of the electrode of interest.
------------------	---------------------------------------

Returns

The SOC of the other electrode that was factored out.

6.10.3.3 to_json() `def ep_bolfi.utility.fitting_functions.OCV_fit_result.to_json (`
 `self)`

6.10.4 Member Data Documentation

6.10.4.1 a `ep_bolfi.utility.fitting_functions.OCV_fit_result.a`

The a (inverse plateau widths) parameters.

6.10.4.2 E₀ `ep_bolfi.utility.fitting_functions.OCV_fit_result.E_0`

The E₀ (plateau voltages) parameters.

6.10.4.3 fit `ep_bolfi.utility.fitting_functions.OCV_fit_result.fit`

The fit parameters of the OCV function from Birkel.

et al., excluding the estimated SOC range.

6.10.4.4 function_string `ep_bolfi.utility.fitting_functions.OCV_fit_result.function_string`

The string representation of the interpolating spline fitted to the inverse.

6.10.4.5 OCV `ep_bolfi.utility.fitting_functions.OCV_fit_result.OCV`

The OCV data points.

May be adjusted from the original data.

6.10.4.6 optimize_result `ep_bolfi.utility.fitting_functions.OCV_fit_result.optimize_result`

The `scipy.optimize.OptimizeResult` that led to the fit.

6.10.4.7 SOC `ep_bolfi.utility.fitting_functions.OCV_fit_result.SOC`

The SOC data points.

6.10.4.8 SOC_offset `ep_bolfi.utility.fitting_functions.OCV_fit_result.SOC_offset`

If another electrode was factored out in the data, this may contain its SOC at SOC 0 of the electrode of interest.

6.10.4.9 SOC_range ep_bolfi.utility.fitting_functions.OCV_fit_result.SOC_range

The SOC range of the data.

6.10.4.10 SOC_scale ep_bolfi.utility.fitting_functions.OCV_fit_result.SOC_scale

If another electrode was factored out in the data, this may contain the the rate of change of its SOC to that of the electrode of interest.

6.10.4.11 spline_interpolation_coefficients ep_bolfi.utility.fitting_functions.OCV_fit_result.spline_interpolation_↔
coefficients

The coefficients of the interpolating spline fitted to the inverse.

6.10.4.12 spline_interpolation_knots ep_bolfi.utility.fitting_functions.OCV_fit_result.spline_interpolation_knots

The knots of the interpolating spline fitted to the inverse.

6.10.4.13 Δx ep_bolfi.utility.fitting_functions.OCV_fit_result.Δx

The Δx (phase proportion) parameters.

The documentation for this class was generated from the following file:

- [utility/fitting_functions.py](#)

6.11 ep_bolfi.optimization.EP_BOLFI.Optimizer_State Class Reference

Handles the heuristics for the EP-BOLFI operation modes.

Public Member Functions

- `def __init__ (self, input_dim, mcmc_chains, total_evidence, posterior_samples, gelman_rubin_threshold, ess_ratio_resample=5.0, ess_ratio_sampling_from_zero=-1.0, ess_ratio_abort=20.0, posterior_sampling_increase=1.↔2, model_resampling_increase=1.2)`
- `def calculate_next_step (self, ess_ratio, action=None)`

Public Attributes

- [input_dim](#)
- [mcmc_chains](#)
- [total_evidence](#)
- [posterior_samples](#)
- [gelman_rubin_threshold](#)
- [ess_ratio_resample](#)
- [ess_ratio_sampling_from_zero](#)
- [ess_ratio_abort](#)
- [posterior_sampling_increase](#)
- [model_resampling_increase](#)
- [finished](#)
- [sampling_from_zero](#)
- [resampling](#)
- [initials](#)
- [order_of_actions](#)
- [current_action](#)
- [verbose_actions](#)

6.11.1 Detailed Description

Handles the heuristics for the EP-BOLFI operation modes.

6.11.2 Constructor & Destructor Documentation

6.11.2.1 `__init__()` `def ep_bolfi.optimization.EP_BOLFI.Optimizer_State.__init__(`
`self,`
`input_dim,`
`mcmc_chains,`
`total_evidence,`
`posterior_samples,`
`gelman_rubin_threshold,`
`ess_ratio_resample = 5.0,`
`ess_ratio_sampling_from_zero = -1.0,`
`ess_ratio_abort = 20.0,`
`posterior_sampling_increase = 1.2,`
`model_resampling_increase = 1.2)`

6.11.3 Member Function Documentation

6.11.3.1 `calculate_next_step()` `def ep_bolfi.optimization.EP_BOLFI.Optimizer_State.calculate_next_step(`
`self,`
`ess_ratio,`
`action = None)`

6.11.4 Member Data Documentation

6.11.4.1 current_action ep_bolfi.optimization.EP_BOLFI.Optimizer_State.current_action

6.11.4.2 ess_ratio_abort ep_bolfi.optimization.EP_BOLFI.Optimizer_State.ess_ratio_abort

6.11.4.3 ess_ratio_resample ep_bolfi.optimization.EP_BOLFI.Optimizer_State.ess_ratio_resample

6.11.4.4 ess_ratio_sampling_from_zero ep_bolfi.optimization.EP_BOLFI.Optimizer_State.ess_ratio_sampling_from_zero

6.11.4.5 finished ep_bolfi.optimization.EP_BOLFI.Optimizer_State.finished

6.11.4.6 gelman_rubin_threshold ep_bolfi.optimization.EP_BOLFI.Optimizer_State.gelman_rubin_threshold

6.11.4.7 initials ep_bolfi.optimization.EP_BOLFI.Optimizer_State.initials

6.11.4.8 input_dim ep_bolfi.optimization.EP_BOLFI.Optimizer_State.input_dim

6.11.4.9 mcmc_chains ep_bolfi.optimization.EP_BOLFI.Optimizer_State.mcmc_chains

6.11.4.10 model_resampling_increase ep_bolfi.optimization.EP_BOLFI.Optimizer_State.model_resampling_increase

6.11.4.11 order_of_actions `ep_bolfi.optimization.EP_BOLFI.Optimizer_State.order_of_actions`

6.11.4.12 posterior_samples `ep_bolfi.optimization.EP_BOLFI.Optimizer_State.posterior_samples`

6.11.4.13 posterior_sampling_increase `ep_bolfi.optimization.EP_BOLFI.Optimizer_State.posterior_sampling_increase`

6.11.4.14 resampling `ep_bolfi.optimization.EP_BOLFI.Optimizer_State.resampling`

6.11.4.15 sampling_from_zero `ep_bolfi.optimization.EP_BOLFI.Optimizer_State.sampling_from_zero`

6.11.4.16 total_evidence `ep_bolfi.optimization.EP_BOLFI.Optimizer_State.total_evidence`

6.11.4.17 verbose_actions `ep_bolfi.optimization.EP_BOLFI.Optimizer_State.verbose_actions`

The documentation for this class was generated from the following file:

- optimization/[EP_BOLFI.py](#)

6.12 `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator` Class Reference

Normalizes sampling to a standard normal distribution.

Public Member Functions

- `def __init__ (self, simulator, fixed_parameters, free_parameters_names, r, Q, experimental_data, feature_extractor, transform_parameters={}, fixed_parameter_order=None)`
- `def search_to_transformed_trial (self, search_space_parameters)`
Transforms search space parameters to model ones.
- `def transformed_trial_to_search (self, model_space_parameters)`
Transforms model space parameters to model ones.
- `def undo_transformation (self, transformed_trial_parameters)`
Undo the transforms in 'self.transform_parameters'.
- `def apply_transformation (self, trial_parameters)`
Apply the transforms in 'self.transform_parameters'.
- `def elfi_simulator (self, *args, **kwargs)`
A battery model simulator that can be used with ELFI.

Public Attributes

- [simulator](#)
- [fixed_parameters](#)
- [free_parameters_names](#)
- [fixed_parameter_order](#)
- [r](#)
- [Q](#)
- [experimental_data](#)
- [feature_extractor](#)
- [transform_parameters](#)
- [log_of_tried_parameters](#)
 - Stores all parameter combinations that have been tried.*
- [experimental_features](#)
 - Extract the features from the experimental data.*
- [input_dim](#)
 - Input dimension of the estimation task.*
- [output_dim](#)
 - Output dimension of the estimation task (number of features).*
- [add_parameters](#)
 - Create a function to combine the free and fixed parameters.*
- [back_transform_matrix](#)
 - Compute the linear transformation of parameters for which the covariance of the underlying multivariate normal distribution is a diagonal matrix.*
- [variances](#)
 - Variances of the model parameters.*
- [transform_matrix](#)
 - Inverse of back_transform_matrix.*
- [transformed_means](#)
 - transform_matrix @ Q @ back_transform_matrix is diagonal.*
- [norm_factor](#)
 - Now that the multivariate normal distribution is decomposed into various univariate ones, norm them to have equal variance 1.*
- [un_norm_factor](#)
 - Inverse of norm_factor.*
- [normed_means](#)
 - Expectation values of the normed univariate normal distributions.*

6.12.1 Detailed Description

Normalizes sampling to a standard normal distribution.

In order to help BOLFI to work efficiently with the least amount of setup required, this class mediates between the model parameters and a standard normal distribution for sampling. In a sense, the simulator output gets transformed into covariance eigenvectors.

6.12.2 Constructor & Destructor Documentation

6.12.2.1 `__init__()` `def ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.__init__(`
`self,`
`simulator,`
`fixed_parameters,`
`free_parameters_names,`
`r,`
`Q,`
`experimental_data,`
`feature_extractor,`
`transform_parameters = {},`
`fixed_parameter_order = None)`

Parameters

<i>simulator</i>	The function that returns results given parameters.
<i>fixed_parameters</i>	Dictionary of parameters that stay fixed and their values.
<i>free_parameters_names</i>	List of the names of parameters which shall be inferred.
<i>r</i>	'Q' times the mean of the distribution of free parameters.
<i>Q</i>	Inverse covariance matrix of free parameters, i.e. precision. It is used to transform the free parameters given to the 'simulator' into the ones used in the model. Most notably, these univariate standard normal distributions get transformed into a multivariate normal distribution corresponding to Q and r.
<i>experimental_data</i>	The experimental data that the model will be fitted to. It has to have the same structure as the 'simulator' output.
<i>feature_extractor</i>	A function that takes the output of 'simulator' / the 'experimental_data' and returns a list of numbers, the features.
<i>transform_parameters</i>	Optional transformations between the parameter space that is used for searching for optimal parameters and the battery model parameters. 'Q' and 'r' define a normal distribution in that search space. The keys are the names of the free parameters. The values are 2-tuples. The first entry is a function taking the search space parameter and returning the model parameter. The second entry is the inverse function.
<i>fixed_parameter_order</i>	Optional fixed parameter order. Prevents erroneous behaviour when the parameter dictionaries get reshuffled. Highly recommended.

6.12.3 Member Function Documentation

6.12.3.1 `apply_transformation()` `def ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.apply_transformation(`
`self,`
`trial_parameters)`

Apply the transforms in 'self.transform_parameters'.

Parameters

<i>trial_parameters</i>	A dictionary. The keys are the 'free_parameters_names' and the values are the actual model parameters.
-------------------------	--

Returns

The given dictionary with the vales transformed to the modified parameter space as specified in 'self.transform_parameters'.

6.12.3.2 elfi_simulator() `def ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.elfi_simulator (self, * args, ** kwargs)`

A battery model simulator that can be used with ELFI.

Parameters

<i>*args</i>	The parameters as given by the prior nodes. Their order has to correspond to that of the parameter 'free_parameters' given to 'return_simulator'.
<i>**kwargs</i>	Keyword parameters batch_size and random_state, but both are unused (they just get passed by BOLFI).

Returns

Simulated features for the given free parameters.

6.12.3.3 search_to_transformed_trial() `def ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.search_to_transformed_trial (self, search_space_parameters)`

Transforms search space parameters to model ones.

Parameters

<i>search_space_parameters</i>	A list of lists which each contain a single search space parameter sample as it is returned by the sample functions of ELFI. In the case of only sample, a list also works.
--------------------------------	---

Returns

A dictionary with its keys as the names of the parameters. Their order in the 'search_space_parameters' is given by the order of 'self.free_parameters_names'. The values yield the model parameters when passed through the functions in 'self.transform_parameters'.

6.12.3.4 transformed_trial_to_search() `def ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.transformed_trial_to_search (`

```
self,
model_space_parameters )
```

Transforms model space parameters to model ones.

Parameters

<i>model_space_parameters</i>	A dictionary. The keys are the 'self.free_parameters_names' and the values are the model parameters after applying the transformations given in 'self.transform_parameters'.
-------------------------------	--

Returns

A list (of lists) which each contain a single search space parameter sample as it is returned by the sample functions of ELFI. If the 'model_space_parameters' dictionary values are numbers, the returned value is a list. If they are lists, the returned value is a list of corresponding lists. In that case, each and every list must have the same length.

6.12.3.5 undo_transformation() `def ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.undo_transformation (`

```
self,
transformed_trial_parameters )
```

Undo the transforms in 'self.transform_parameters'.

Parameters

<i>transformed_trial_parameters</i>	A dictionary. The keys are the 'free_parameters_names' and the values are the model parameters after they have been transformed as specified in 'self.transform_parameters'.
-------------------------------------	--

Returns

The given dictionary with the values transformed back to the actual model parameter values.

6.12.4 Member Data Documentation

6.12.4.1 add_parameters `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.add_parameters`

Create a function to combine the free and fixed parameters.

6.12.4.2 back_transform_matrix `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.back_transform_matrix`

Compute the linear transformation of parameters for which the covariance of the underlying multivariate normal distribution is a diagonal matrix.

That is, compute the eigenvectors of Q. It is more stable since Q has growing eigenvectors in convergence.

6.12.4.3 `experimental_data` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.experimental_data`

6.12.4.4 `experimental_features` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.experimental_features`

Extract the features from the experimental data.

6.12.4.5 `feature_extractor` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.feature_extractor`

6.12.4.6 `fixed_parameter_order` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.fixed_parameter_order`

6.12.4.7 `fixed_parameters` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.fixed_parameters`

6.12.4.8 `free_parameters_names` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.free_parameters_names`

6.12.4.9 `input_dim` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.input_dim`

Input dimension of the estimation task.

6.12.4.10 `log_of_tried_parameters` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.log_of_tried_parameters`

Stores all parameter combinations that have been tried.

6.12.4.11 `norm_factor` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.norm_factor`

Now that the multivariate normal distribution is decomposed into various univariate ones, norm them to have equal variance 1.

6.12.4.12 `normed_means` `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.normed_means`

Expectation values of the normed univariate normal distributions.

6.12.4.13 output_dim ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.output_dim

Output dimension of the estimation task (number of features).

6.12.4.14 Q ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.Q

6.12.4.15 r ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.r

6.12.4.16 simulator ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.simulator

6.12.4.17 transform_matrix ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.transform_matrix

Inverse of back_transform_matrix.

6.12.4.18 transform_parameters ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.transform_parameters

6.12.4.19 transformed_means ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.transformed_means

transform_matrix @ Q @ back_transform_matrix is diagonal.

The correct transformation for vectors v is then transform_matrix @ v . The product below corresponds to Q^{-1} @ r . It is just expressed in the eigenvector space of Q for efficiency.

6.12.4.20 un_norm_factor ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.un_norm_factor

Inverse of norm_factor.

6.12.4.21 variances ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator.variances

Variances of the model parameters.

The documentation for this class was generated from the following file:

- optimization/[EP_BOLFI.py](#)

6.13 ep_bolfi.utility.dataset_formatting.Static_Information Class Reference

Contains additional informations, e.g.

Inheritance diagram for ep_bolfi.utility.dataset_formatting.Static_Information:

Collaboration diagram for ep_bolfi.utility.dataset_formatting.Static_Information:

Public Member Functions

- def `__init__` (self, [timepoints](#), [currents](#), [voltages](#), [other_columns](#)={}, [indices](#)=None)
- def `to_json` (self)
- def `subslice` (self, start, stop, step=1)
Selects a subslice of the data segments.
- def `subarray` (self, array)
Selects the data segments with the given indices.
- def `extend` (self, other)

Public Attributes

- [asymptotic_voltages](#)
The voltages that the voltage curve seems to converge to in a segment.
- [ir_steps](#)
The instantaneous IR drops before each segment.
- [exp_I_decays](#)
Same as `exp_U_decays` for current decays (PITT).
- [exp_U_decays](#)
The fit parameters of the exponential voltage decays in each segment.

6.13.1 Detailed Description

Contains additional informations, e.g.

for GITT. Each member variable is a list and has the same length as the other ones.

6.13.2 Constructor & Destructor Documentation

6.13.2.1 `__init__()` def ep_bolfi.utility.dataset_formatting.Static_Information.__init__ (self, timepoints, currents, voltages, other_columns = {}, indices = None)

Reimplemented from [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.13.3 Member Function Documentation

6.13.3.1 extend() `def ep_bolfi.utility.dataset_formatting.Static_Information.extend (`
 self,
 other)

Reimplemented from [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.13.3.2 subarray() `def ep_bolfi.utility.dataset_formatting.Static_Information.subarray (`
 self,
 array)

Selects the data segments with the given indices.

Parameters

<i>array</i>	The indices of the segments that are to be returned.
--------------	--

Returns

A new [Static_Information](#) object containing the subset.

Reimplemented from [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.13.3.3 subslice() `def ep_bolfi.utility.dataset_formatting.Static_Information.subslice (`
 self,
 start,
 stop,
 step = 1)

Selects a subslice of the data segments.

The arguments exactly match the `slice(...)` notation.

Parameters

<i>start</i>	The index of the first segment to be included.
<i>stop</i>	The index of the first segment to not be included.
<i>step</i>	Steps between selected segments. Default: 1.

Returns

A new [Static_Information](#) object containing the slice.

Reimplemented from [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.13.3.4 `to_json()` `def ep_bolfi.utility.dataset_formatting.Static_Information.to_json (self)`

Reimplemented from [ep_bolfi.utility.dataset_formatting.Cycling_Information](#).

6.13.4 Member Data Documentation

6.13.4.1 `asymptotic_voltages` `ep_bolfi.utility.dataset_formatting.Static_Information.asymptotic_voltages`

The voltages that the voltage curve seems to converge to in a segment.

Only makes sense for those segments that are rest periods or when the OCV was subtracted.

6.13.4.2 `exp_I_decays` `ep_bolfi.utility.dataset_formatting.Static_Information.exp_I_decays`

Same as `exp_U_decays` for current decays (PITT).

6.13.4.3 `exp_U_decays` `ep_bolfi.utility.dataset_formatting.Static_Information.exp_U_decays`

The fit parameters of the exponential voltage decays in each segment.

Each set of fit parameters is a 3-tuple (a,b,c) where the fit function has the following form: $a + b * \exp(-c * (t - t_end_of_segment))$. Failed or missing fits are best indicated by (NaN, NaN, NaN). """

6.13.4.4 `ir_steps` `ep_bolfi.utility.dataset_formatting.Static_Information.ir_steps`

The instantaneous IR drops before each segment.

Positive values are voltage rises and negative values voltage drops.

The documentation for this class was generated from the following file:

- [utility/dataset_formatting.py](#)

6.14 `ep_bolfi.utility.preprocessing.SubstitutionDict` Class Reference

A dictionary with some automatic substitutions.

Inheritance diagram for `ep_bolfi.utility.preprocessing.SubstitutionDict`:

Collaboration diagram for `ep_bolfi.utility.preprocessing.SubstitutionDict`:

Public Member Functions

- `def __init__ (self, storage, substitutions={})`
- `def __delitem__ (self, key)`
- `def __getitem__ (self, key)`
- `def __iter__ (self)`
- `def __len__ (self)`
- `def __setitem__ (self, key, value)`
- `def __str__ (self)`
- `def __repr__ (self)`
- `def log_lock (self)`
- `def dependent_variables (self, parameters)`

6.14.1 Detailed Description

A dictionary with some automatic substitutions.

"substitutions" is a dictionary that extends "storage" with automatic substitution rules depending on its value types:

- string, which serves the value of "storage" at that value.
- callable which takes one parameter, which will get passed its [SubstitutionDict](#) instance and serves its return value.
- any other type, which serves the value as-is. Assigning values to keys afterwards will overwrite substitutions.

6.14.2 Constructor & Destructor Documentation

6.14.2.1 `__init__()` `def ep_bolfi.utility.preprocessing.SubstitutionDict.__init__(
self,
storage,
substitutions = {})`

6.14.3 Member Function Documentation

6.14.3.1 `__delitem__()` `def ep_bolfi.utility.preprocessing.SubstitutionDict.__delitem__(
self,
key)`

6.14.3.2 `__getitem__()` def ep_bolfi.utility.preprocessing.SubstitutionDict.__getitem__ (
 self,
 key)

6.14.3.3 `__iter__()` def ep_bolfi.utility.preprocessing.SubstitutionDict.__iter__ (
 self)

6.14.3.4 `__len__()` def ep_bolfi.utility.preprocessing.SubstitutionDict.__len__ (
 self)

6.14.3.5 `__repr__()` def ep_bolfi.utility.preprocessing.SubstitutionDict.__repr__ (
 self)

6.14.3.6 `__setitem__()` def ep_bolfi.utility.preprocessing.SubstitutionDict.__setitem__ (
 self,
 key,
 value)

6.14.3.7 `__str__()` def ep_bolfi.utility.preprocessing.SubstitutionDict.__str__ (
 self)

6.14.3.8 `dependent_variables()` def ep_bolfi.utility.preprocessing.SubstitutionDict.dependent_variables (
 self,
 parameters)

6.14.3.9 `log_lock()` def ep_bolfi.utility.preprocessing.SubstitutionDict.log_lock (
 self)

The documentation for this class was generated from the following file:

- [utility/preprocessing.py](#)

7 File Documentation

7.1 `__init__.py` File Reference

Namespaces

- [ep_bolfi](#)

7.2 `models/__init__.py` File Reference

Namespaces

- [ep_bolfi.models](#)

7.3 `optimization/__init__.py` File Reference

Namespaces

- [ep_bolfi.optimization](#)

7.4 `utility/__init__.py` File Reference

Namespaces

- [ep_bolfi.utility](#)

7.5 `models/assess_effective_parameters.py` File Reference

Classes

- class [ep_bolfi.models.assess_effective_parameters.Effective_Parameters](#)
Calculates, stores and prints effective parameters.

Namespaces

- [ep_bolfi.models.assess_effective_parameters](#)
Evaluates a parameter set for, e.g., overpotential and capacity.

7.6 `models/electrolyte.py` File Reference

Classes

- class [ep_bolfi.models.electrolyte.Electrolyte_internal](#)
Defining equations for a symmetric Li cell with electrolyte.
- class [ep_bolfi.models.electrolyte.Electrolyte](#)
Electrolyte model assuming a symmetric Li-metal cell.

Namespaces

- [ep_bolfi.models.electrolyte](#)

Contains a PyBaMM-compatible electrolyte model.

7.7 models/solversetup.py File Reference

Namespaces

- [ep_bolfi.models.solversetup](#)

This file eases the setup and simulation of PyBaMM battery models.

Functions

- def [ep_bolfi.models.solversetup.solver_setup](#) (model, parameters, submesh_types, var_pts, spatial_↵ methods, geometry=None, reltol=1e-6, abstol=1e-6, root_tol=1e-3, dt_max=None, free_parameters=[], verbose=False, logging_file=None)
Processes the model and returns a runnable solver.
- def [ep_bolfi.models.solversetup.simulation_setup](#) (model, operation_input, parameters, submesh_types, var_pts, spatial_methods, geometry=None, reltol=1e-6, abstol=1e-6, root_tol=1e-3, dt_max=None, free_↵ parameters=[], verbose=False, logging_file=None)
Processes the model and returns a runnable solver.
- def [ep_bolfi.models.solversetup.auto_var_pts](#) (x_n, x_s, x_p, r_n, r_p, y=1, z=1)
Utility function for setting the discretization density.
- def [ep_bolfi.models.solversetup.spectral_mesh_pts_and_method](#) (order_s_n, order_s_p, order_↵ e, volumes_e_n=1, volumes_e_s=1, volumes_e_p=1, halfcell=False)
Utility function for default mesh and spatial methods.

7.8 models/standard_parameters.py File Reference

Namespaces

- [ep_bolfi.models.standard_parameters](#)

Comprehensive list of parameters of every model.

Functions

- def [ep_bolfi.models.standard_parameters.SOC_n_dim_init](#) (x)
Initial SOC of the anode.
- def [ep_bolfi.models.standard_parameters.SOC_n_init](#) (x)
Non-dimensionalized initial SOC of the anode.
- def [ep_bolfi.models.standard_parameters.SOC_p_dim_init](#) (x)
Initial SOC of the cathode.
- def [ep_bolfi.models.standard_parameters.SOC_p_init](#) (x)
Non-dimensionalized initial SOC of the cathode.
- def [ep_bolfi.models.standard_parameters.De_dim](#) (c_e_dim, T_dim)
Electrolyte diffusivity.
- def [ep_bolfi.models.standard_parameters.De](#) (c_e, T)
Non-dimensionalized electrolyte diffusivity.

- def `ep_bolfi.models.standard_parameters.k_e_dim` (c_e_dim , T_dim)
Electrolyte conductivity.
- def `ep_bolfi.models.standard_parameters.k_e` (c_e , T)
Non-dimensionalized electrolyte conductivity.
- def `ep_bolfi.models.standard_parameters.t_plus_dim` (c_e_dim)
Transference number.
- def `ep_bolfi.models.standard_parameters.t_plus` (c_e)
Non-dimensionalized (referring to the input) transference number.
- def `ep_bolfi.models.standard_parameters.one_plus_dlnf_dlnc_dim` (c_e_dim)
Thermodynamic factor.
- def `ep_bolfi.models.standard_parameters.one_plus_dlnf_dlnc` (c_e)
Non-dimensionalized (referring to the input) thermodynamic factor.
- def `ep_bolfi.models.standard_parameters.D_n_dim` (SOC_n , T_dim)
Anode diffusivity.
- def `ep_bolfi.models.standard_parameters.D_n` (SOC_n , T)
Non-dimensionalized anode diffusivity.
- def `ep_bolfi.models.standard_parameters.D_p_dim` (SOC_p , T_dim)
Cathode diffusivity.
- def `ep_bolfi.models.standard_parameters.D_p` (SOC_p , T)
Non-dimensionalized cathode diffusivity.
- def `ep_bolfi.models.standard_parameters.i_sn_0_dim` (c_n_dim , $SOC_n_surf_dim$, c_n_max , T_dim)
Anode exchange current density.
- def `ep_bolfi.models.standard_parameters.i_sn_0` (c_n , SOC_n_surf , c_n_max , T)
Non-dimensionalized anode exchange current density.
- def `ep_bolfi.models.standard_parameters.d_c_n_i_sn_0_dim` (c_n_dim , $SOC_n_surf_dim$, c_n_max , T_dim)
 ∂ anode exchange current density / ∂ electrolyte concentration.
- def `ep_bolfi.models.standard_parameters.d_c_n_i_sn_0` (c_n , SOC_n_surf , c_n_max , T)
The non-dimensionalized version of the prior variable.
- def `ep_bolfi.models.standard_parameters.i_sep_0_dim` (c_{ep_dim} , $SOC_p_surf_dim$, c_p_max , T_dim)
Cathode exchange current density.
- def `ep_bolfi.models.standard_parameters.i_sep_0` (c_{ep} , SOC_p_surf , c_p_max , T)
Non-dimensionalized cathode exchange current density.
- def `ep_bolfi.models.standard_parameters.d_c_ep_i_sep_0_dim` (c_{ep_dim} , $SOC_p_surf_dim$, c_p_max , T_dim)
 ∂ cathode exchange current density / ∂ electrolyte concentration.
- def `ep_bolfi.models.standard_parameters.d_c_ep_i_sep_0` (c_{ep} , SOC_p_surf , c_p_max , T)
The non-dimensionalized version of the prior variable.
- def `ep_bolfi.models.standard_parameters.dOCV_n_dT_dim` (SOC_n)
 ∂ anode OCV / ∂ temperature.
- def `ep_bolfi.models.standard_parameters.dOCV_n_dT` (SOC_n)
Non-dimensionalized ∂ anode OCV / ∂ temperature.
- def `ep_bolfi.models.standard_parameters.dOCV_n_dT_dSOC_n_dim` (SOC_n)
 $(\partial$ anode OCV / ∂ temperature) / ∂ anode SOC.
- def `ep_bolfi.models.standard_parameters.dOCV_n_dT_dSOC_n` (SOC_n)
Non-dimensionalized $(\partial$ anode OCV / ∂ temperature) / ∂ anode SOC.
- def `ep_bolfi.models.standard_parameters.dOCV_p_dT_dim` (SOC_p)
 ∂ cathode OCV / ∂ temperature.
- def `ep_bolfi.models.standard_parameters.dOCV_p_dT` (SOC_p)
Non-dimensionalized ∂ cathode OCV / ∂ temperature.
- def `ep_bolfi.models.standard_parameters.dOCV_p_dT_dSOC_p_dim` (SOC_p)
 $(\partial$ cathode OCV / ∂ temperature) / ∂ cathode SOC.
- def `ep_bolfi.models.standard_parameters.dOCV_p_dT_dSOC_p` (SOC_p)

- Non-dimensionalized (∂ cathode OCV / ∂ temperature) / ∂ cathode SOC.*
- def `ep_bolfi.models.standard_parameters.OCV_n_dim` (SOC_n, T_{dim})
Anode OCV.
- def `ep_bolfi.models.standard_parameters.OCV_n` (SOC_n, T)
Non-dimensionalized anode OCV.
- def `ep_bolfi.models.standard_parameters.dOCV_n_dim_dSOC_n` (SOC_n, T_{dim})
 ∂ anode OCV / ∂ anode SOC.
- def `ep_bolfi.models.standard_parameters.dOCV_n_dSOC_n` (SOC_n, T)
Non-dimensionalized ∂ anode OCV / ∂ anode SOC.
- def `ep_bolfi.models.standard_parameters.OCV_p_dim` (SOC_p, T_{dim})
Cathode OCV.
- def `ep_bolfi.models.standard_parameters.OCV_p` (SOC_p, T)
Non-dimensionalized cathode OCV.
- def `ep_bolfi.models.standard_parameters.dOCV_p_dim_dSOC_p` (SOC_p, T_{dim})
 ∂ cathode OCV / ∂ cathode SOC.
- def `ep_bolfi.models.standard_parameters.dOCV_p_dSOC_p` (SOC_p, T)
Non-dimensionalized ∂ cathode OCV / ∂ cathode SOC.

Variables

- `ep_bolfi.models.standard_parameters.R` = `Scalar(constants.R)`
- `ep_bolfi.models.standard_parameters.F` = `Scalar(constants.physical_constants["Faraday constant"])[0]`
- `ep_bolfi.models.standard_parameters.k_B` = `constants.physical_constants["Boltzmann constant"]`
- `ep_bolfi.models.standard_parameters.q_e` = `constants.physical_constants["electron volt"]`
- `ep_bolfi.models.standard_parameters.T_ref` = `Parameter("Reference temperature [K]")`
- `ep_bolfi.models.standard_parameters.T_init` = `Parameter("Initial temperature [K]")`
- `ep_bolfi.models.standard_parameters.thermal_voltage` = `R * T_ref / F`
- `ep_bolfi.models.standard_parameters.ΔT` = `Scalar(1)`
- `ep_bolfi.models.standard_parameters.L_n_dim` = `Parameter("Negative electrode thickness [m]")`
- `ep_bolfi.models.standard_parameters.L_s_dim` = `Parameter("Separator thickness [m]")`
- `ep_bolfi.models.standard_parameters.L_p_dim` = `Parameter("Positive electrode thickness [m]")`
- `ep_bolfi.models.standard_parameters.L_dim` = `L_n_dim + L_s_dim + L_p_dim`
- `ep_bolfi.models.standard_parameters.A` = `Parameter("Current collector perpendicular area [m2]")`
- `ep_bolfi.models.standard_parameters.V` = `Parameter("Cell volume [m3]")`
- `ep_bolfi.models.standard_parameters.L_x` = `L_dim`
- `ep_bolfi.models.standard_parameters.L_y` = `Parameter("Electrode width [m]")`
- `ep_bolfi.models.standard_parameters.L_z` = `Parameter("Electrode height [m]")`
- `ep_bolfi.models.standard_parameters.C` = `Parameter("Typical current [A]")`
- `ep_bolfi.models.standard_parameters.U_l` = `pybamm.Parameter("Lower voltage cut-off [V]")`
- `ep_bolfi.models.standard_parameters.U_u` = `pybamm.Parameter("Upper voltage cut-off [V]")`
- `ep_bolfi.models.standard_parameters.c_e_typ` = `pybamm.Parameter("Typical electrolyte concentration [mol.m-3]")`
- `ep_bolfi.models.standard_parameters.c_n` = `pybamm.Parameter("Maximum concentration in negative electrode [mol.m-3]")`
- `ep_bolfi.models.standard_parameters.c_p` = `pybamm.Parameter("Maximum concentration in positive electrode [mol.m-3]")`
- `ep_bolfi.models.standard_parameters.σ_n_dim` = `pybamm.Parameter("Negative electrode conductivity [S.m-1]")`
- `ep_bolfi.models.standard_parameters.σ_p_dim` = `pybamm.Parameter("Positive electrode conductivity [S.m-1]")`
- `ep_bolfi.models.standard_parameters.a_n_dim` = `Parameter("Negative electrode surface area to volume ratio [m-1]")`

- `ep_bolfi.models.standard_parameters.a_p_dim` = Parameter("Positive electrode surface area to volume ratio [m⁻¹]")
- `ep_bolfi.models.standard_parameters.R_n` = Parameter("Negative particle radius [m]")
- `ep_bolfi.models.standard_parameters.R_p` = Parameter("Positive particle radius [m]")
- `ep_bolfi.models.standard_parameters.αnn` = Parameter("Negative electrode anodic charge-transfer coefficient")
- `ep_bolfi.models.standard_parameters.αpn` = Parameter("Negative electrode cathodic charge-transfer coefficient")
- `ep_bolfi.models.standard_parameters.αnp` = Parameter("Positive electrode anodic charge-transfer coefficient")
- `ep_bolfi.models.standard_parameters.αpp` = Parameter("Positive electrode cathodic charge-transfer coefficient")
- `ep_bolfi.models.standard_parameters.βa_scalar`
- `ep_bolfi.models.standard_parameters.βes_scalar` = Parameter("Separator Bruggeman coefficient (electrolyte)")
- `ep_bolfi.models.standard_parameters.βep_scalar`
- `ep_bolfi.models.standard_parameters.βa` = pybamm.PrimaryBroadcast(β_a_scalar, "negative electrode")
- `ep_bolfi.models.standard_parameters.βes` = pybamm.PrimaryBroadcast(β_{es}_scalar, "separator")
- `ep_bolfi.models.standard_parameters.βep` = pybamm.PrimaryBroadcast(β_{ep}_scalar, "positive electrode")
- `ep_bolfi.models.standard_parameters.βsn_scalar` = Parameter("Negative electrode Bruggeman coefficient (electrode)")
- `ep_bolfi.models.standard_parameters.βss_scalar` = Parameter("Separator Bruggeman coefficient (electrode)")
- `ep_bolfi.models.standard_parameters.βsp_scalar` = Parameter("Positive electrode Bruggeman coefficient (electrode)")
- `ep_bolfi.models.standard_parameters.βe` = pybamm.Concatenation(β_a, β_{es}, β_{ep})
- `ep_bolfi.models.standard_parameters.εn_scalar` = Parameter("Negative electrode porosity")
- `ep_bolfi.models.standard_parameters.εs_scalar` = Parameter("Separator porosity")
- `ep_bolfi.models.standard_parameters.εp_scalar` = Parameter("Positive electrode porosity")
- `ep_bolfi.models.standard_parameters.εn` = pybamm.PrimaryBroadcast(ε_n_scalar, "negative electrode")
- `ep_bolfi.models.standard_parameters.εs` = pybamm.PrimaryBroadcast(ε_s_scalar, "separator")
- `ep_bolfi.models.standard_parameters.εp` = pybamm.PrimaryBroadcast(ε_p_scalar, "positive electrode")
- `ep_bolfi.models.standard_parameters.ε` = pybamm.Concatenation(ε_n, ε_s, ε_p)
- `ep_bolfi.models.standard_parameters.ε0` = ε * β_e
- `ep_bolfi.models.standard_parameters.zn` = Parameter("Negative electrode electrons in reaction")
- `ep_bolfi.models.standard_parameters.zp` = Parameter("Positive electrode electrons in reaction")
- `ep_bolfi.models.standard_parameters.ce_dim_init` = Parameter("Initial concentration in electrolyte [mol.↔m⁻³]")
- `ep_bolfi.models.standard_parameters.ce_init` = c_e_dim_init / c_e_typ
- `def ep_bolfi.models.standard_parameters.De_typ` = D_e_dim(c_e_typ, T_{ref})
- `def ep_bolfi.models.standard_parameters.κe_typ` = κ_e_dim(c_e_typ, T_{ref})
- `tuple ep_bolfi.models.standard_parameters.κe_hat` = (R * T_{ref} / F) / (C / A * L_{dim} / κ_e_typ)
- `def ep_bolfi.models.standard_parameters.Dn_typ` = D_n_dim(Scalar(0.5), T_{ref})
- `def ep_bolfi.models.standard_parameters.Dp_typ` = D_p_dim(Scalar(0.5), T_{ref})
- `def ep_bolfi.models.standard_parameters.isn_0_ref` = i_{sn}_0_dim(c_e_typ, 0.5 * c_n, c_n, T_{ref})
- `def ep_bolfi.models.standard_parameters.isep_0_ref` = i_{sep}_0_dim(c_e_typ, 0.5 * c_p, c_p, T_{ref})
- `def ep_bolfi.models.standard_parameters.OCVn_ref` = OCV_n_dim(SOC_n_init(0), T_{ref})
- `def ep_bolfi.models.standard_parameters.OCVp_ref` = OCV_p_dim(SOC_p_init(1), T_{ref})
- `ep_bolfi.models.standard_parameters.an` = a_n_dim * R_n
- `ep_bolfi.models.standard_parameters.ap` = a_p_dim * R_p
- `tuple ep_bolfi.models.standard_parameters.Q` = (1 - ε_p_scalar) * L_p_dim * c_p * z_p * F * A
- `ep_bolfi.models.standard_parameters.τd` = F * c_p * L_{dim} / (C / A)
- `int ep_bolfi.models.standard_parameters.τe` = L_{dim}**2 / D_e_typ
- `int ep_bolfi.models.standard_parameters.τn` = R_n**2 / D_n_typ
- `int ep_bolfi.models.standard_parameters.τp` = R_p**2 / D_p_typ

- `ep_bolfi.models.standard_parameters.τrn` = $F * c_n / (i_{sn_0_ref} * a_n_dim)$
- `ep_bolfi.models.standard_parameters.τrp` = $F * c_p / (i_{sep_0_ref} * a_p_dim)$
- `ep_bolfi.models.standard_parameters.timescale` = τ^d
- `int ep_bolfi.models.standard_parameters.Ce` = τ_e / τ^d
- `int ep_bolfi.models.standard_parameters.Cn` = τ_n / τ^d
- `int ep_bolfi.models.standard_parameters.Cp` = τ_p / τ^d
- `ep_bolfi.models.standard_parameters.Crn` = τ_{rn} / τ^d
- `ep_bolfi.models.standard_parameters.Crp` = τ_{rp} / τ^d
- `ep_bolfi.models.standard_parameters.γe` = c_{e_typ} / c_p
- `ep_bolfi.models.standard_parameters.γn` = c_n / c_p
- `ep_bolfi.models.standard_parameters.γp` = c_p / c_p
- `ep_bolfi.models.standard_parameters.Ln` = L_n_dim / L_dim
- `ep_bolfi.models.standard_parameters.Ls` = L_s_dim / L_dim
- `ep_bolfi.models.standard_parameters.Lp` = L_p_dim / L_dim
- `ep_bolfi.models.standard_parameters.Le`
- `tuple ep_bolfi.models.standard_parameters.σn` = $(thermal_voltage / (C / A * L_dim)) * \sigma_n_dim$
- `tuple ep_bolfi.models.standard_parameters.σp` = $(thermal_voltage / (C / A * L_dim)) * \sigma_p_dim$
- `ep_bolfi.models.standard_parameters.Iextern_dim`
- `ep_bolfi.models.standard_parameters.Iextern` = I_extern_dim / C
- `ep_bolfi.models.standard_parameters.nelectrodes_parallel`
- `ep_bolfi.models.standard_parameters.ncells` = `Parameter("Number of cells connected in series to make a battery")`
- `ep_bolfi.models.standard_parameters.Ityp` = C
- `ep_bolfi.models.standard_parameters.Acc` = A
- `ep_bolfi.models.standard_parameters.current_with_time` = I_extern
- `ep_bolfi.models.standard_parameters.dimensional_current_with_time` = I_extern_dim
- `ep_bolfi.models.standard_parameters.dimensional_current_density_with_time` = I_extern_dim / A
- `ep_bolfi.models.standard_parameters.voltage_low_cut` = U_l
- `ep_bolfi.models.standard_parameters.voltage_high_cut` = U_u
- `ep_bolfi.models.standard_parameters.capacity` = `Parameter("Nominal cell capacity [A.h]")`

7.9 optimization/EP_BOLFI.py File Reference

Classes

- class `ep_bolfi.optimization.EP_BOLFI.NDArrayEncoder`
- class `ep_bolfi.optimization.EP_BOLFI.Preprocessed_Simulator`
Normalizes sampling to a standard normal distribution.
- class `ep_bolfi.optimization.EP_BOLFI.Optimizer_State`
Handles the heuristics for the EP-BOLFI operation modes.
- class `ep_bolfi.optimization.EP_BOLFI.EP_BOLFI`
Expectation Propagation and Bayesian Optimization.

Namespaces

- `ep_bolfi.optimization.EP_BOLFI`
- `ep_bolfi.EP_BOLFI.EP_BOLFI`

This file contains functions to perform Expectation Propagation on simulator models using BOLFI (Bayesian Optimization).

Functions

- def [ep_bolfi.optimization.EP_BOLFI.combine_parameters_to_try](#) (parameters, parameters_to_try_dict)
Give every combination as full parameter sets.
- def [ep_bolfi.optimization.EP_BOLFI.fix_parameters](#) (parameters_to_be_fixed)
Returns a function which sets some parameters in advance.

7.10 utility/dataset_formatting.py File Reference

Classes

- class [ep_bolfi.utility.dataset_formatting.Measurement](#)
Defines common methods for measurement objects.
- class [ep_bolfi.utility.dataset_formatting.Cycling_Information](#)
Contains basic cycling informations.
- class [ep_bolfi.utility.dataset_formatting.Static_Information](#)
Contains additional informations, e.g.
- class [ep_bolfi.utility.dataset_formatting.Impedance_Measurement](#)
Contains basic impedance data.

Namespaces

- [ep_bolfi.utility.dataset_formatting](#)
Defines datatypes for further processing.

Functions

- def [ep_bolfi.utility.dataset_formatting.read_csv_from_measurement_system](#) (path, encoding, number_of_comment_lines, headers, delimiter='t', decimal='.', datatype="cycling", segment_column=-1, segments_to_process=None, current_sign_correction={}, correction_column=-1, flip_voltage_sign=False, flip_imaginary_impedance_sign=False, max_number_of_lines=-1)
Read the measurements as returned by common instruments.
- def [ep_bolfi.utility.dataset_formatting.print_hdf5_structure](#) (h5py_object, depth=1, table_limit=4, verbose_limit=64)
brief Simple HDF5 structure viewer.
- def [ep_bolfi.utility.dataset_formatting.convert_none_notation_to_slicing](#) (h5py_object, index)
Access a slice of an HDF5 object by transferable notation.
- def [ep_bolfi.utility.dataset_formatting.get_hdf5_dataset_by_path](#) (h5py_object, path)
Follow the structure of a HDF object to get a certain part.
- def [ep_bolfi.utility.dataset_formatting.read_hdf5_table](#) (path, data_location, headers, datatype="cycling", segment_location=None, segments_to_process=None, current_sign_correction={}, correction_location=None, flip_voltage_sign=False, flip_imaginary_impedance_sign=False)
Read the measurements as stored in a HDF5 file.

7.11 utility/fitting_functions.py File Reference

Classes

- class [ep_bolfi.utility.fitting_functions.NDArrayEncoder](#)
- class [ep_bolfi.utility.fitting_functions.OCV_fit_result](#)
Contains OCV fit parameters and related information.

Namespaces

- [ep_bolfi.utility.fitting_functions](#)

Various helper and fitting functions for processing measurement curves.

Functions

- def [ep_bolfi.utility.fitting_functions.find_occurrences](#) (sequence, value)
Gives indices in sequence where it is closest to value.
- def [ep_bolfi.utility.fitting_functions.smooth_fit](#) (x, y, order=3, splits=None, w=None, s=None, display=False, derivatives=0)
Calculates a smoothed spline with derivatives.
- def [ep_bolfi.utility.fitting_functions.fit_exponential_decay](#) (timepoints, voltages, recursive_depth=1, threshold=0.95)
Extracts a set amount of exponential decay curves.
- def [ep_bolfi.utility.fitting_functions.fit_sqrt](#) (timepoints, voltages, threshold=0.95)
Extracts a square root at the beginning of the data.
- def [ep_bolfi.utility.fitting_functions.fit_drt](#) (frequencies, impedances, lambda_value=-2.0)
- def [ep_bolfi.utility.fitting_functions.laplace_transform](#) (x, y, s)
Performs a basic laplace transformation.
- def [ep_bolfi.utility.fitting_functions.a_fit](#) (yUminus1)
Calculates the conversion from "A parametric OCV model".
- def [ep_bolfi.utility.fitting_functions.OCV_fit_function](#) (E_OCV, *args, z=1.0, T=298.15, individual=False, fit_SOC_range=False, rescale=False)
The OCV model from "A parametric OCV model".
- def [ep_bolfi.utility.fitting_functions.d_dE_OCV_fit_function](#) (E_OCV, *args, z=1.0, T=298.15, individual=False, fit_SOC_range=False, rescale=False)
The derivative of [fitting_functions.OCV_fit_function](#).
- def [ep_bolfi.utility.fitting_functions.d2_dE2_OCV_fit_function](#) (E_OCV, *args, z=1.0, T=298.15, individual=False, fit_SOC_range=False, rescale=False)
The 2nd derivative of [fitting_functions.OCV_fit_function](#).
- def [ep_bolfi.utility.fitting_functions.inverse_OCV_fit_function](#) (SOC, *args, z=1.0, T=298.15, inverted=True)
The inverse of [fitting_functions.OCV_fit_function](#).
- def [ep_bolfi.utility.fitting_functions.inverse_d_dSOC_OCV_fit_function](#) (SOC, *args, z=1.0, T=298.15, inverted=True)
The derivative of the inverse of [fitting_functions.OCV_fit_function](#).
- def [ep_bolfi.utility.fitting_functions.inverse_d2_dSOC2_OCV_fit_function](#) (SOC, *args, z=1.0, T=298.15, inverted=True)
The 2nd derivative of the inverse of [fitting_functions.OCV_fit_function](#).
- def [ep_bolfi.utility.fitting_functions.fit_OCV](#) (SOC, OCV, N=4, SOC_range_bounds=(0.2, 0.8), SOC_range_limits=(0.0, 1.0), z=1.0, T=298.15, inverted=True, fit_SOC_range=True, distance_order=2, weights=None, initial_parameters=None, minimize_options=None)
Fits data to [fitting_functions.OCV_fit_function](#).
- def [ep_bolfi.utility.fitting_functions.verbose_spline_parameterization](#) (coeffs, knots, order, format='python', function_name="OCV", function_args="SOC", derivatives=0, spline_transformation="", verbose=False)
Gives the monomic representation of a B-spline.

7.12 utility/preprocessing.py File Reference

Classes

- class [ep_bolfi.utility.preprocessing.SubstitutionDict](#)
A dictionary with some automatic substitutions.

Namespaces

- [ep_bolfi.utility.preprocessing](#)

Contains frequently used workflows in dataset preprocessing.

Functions

- [def ep_bolfi.utility.preprocessing.fix_parameters](#) (parameters_to_be_fixed)
Returns a function which sets some parameters in advance.
- [def ep_bolfi.utility.preprocessing.combine_parameters_to_try](#) (parameters, parameters_to_try_dict)
Give every combination as full parameter sets.
- [def ep_bolfi.utility.preprocessing.calculate_means_and_standard_deviations](#) (mean, covariance, free_↵ parameters_names, transform_parameters={}, bounds_in_standard_deviations=1, **kwargs)
Calculate means and standard deviations.
- [def ep_bolfi.utility.preprocessing.approximate_confidence_ellipsoid](#) (parameters, free_parameters_names, covariance, mean=None, transform_parameters={}, refinement=True, confidence=0.95)
Approximate a confidence ellipsoid.
- [def ep_bolfi.utility.preprocessing.capacity](#) (parameters, electrode="positive")
Convenience function for calculating the capacity.
- [def ep_bolfi.utility.preprocessing.calculate_SOC](#) (timepoints, currents, initial_SOC=0, sign=1, capacity=1)
Transforms applied current over time into SOC.
- [def ep_bolfi.utility.preprocessing.calculate_both_SOC_from_OCV](#) (parameters, negative_SOC_from_↵ cell_SOC, positive_SOC_from_cell_SOC, OCV)
Calculates the SOC of both electrodes from their OCV.
- [def ep_bolfi.utility.preprocessing.subtract_OCV_curve_from_cycles](#) (dataset, parameters, starting_↵ SOC=None, starting_OCV=None, electrode="positive", current_sign=0, voltage_sign=0)
Removes the OCV curve from a cycling measurement.
- [def ep_bolfi.utility.preprocessing.subtract_both_OCV_curves_from_cycles](#) (dataset, parameters, negative_↵ _SOC_from_cell_SOC, positive_SOC_from_cell_SOC, starting_SOC=None, starting_OCV=None)
Removes the OCV curve from a single cycle.
- [def ep_bolfi.utility.preprocessing.laplace_transform](#) (x, y, s)
Performs a basic laplace transformation.
- [def ep_bolfi.utility.preprocessing.find_occurrences](#) (sequence, value)
Gives indices in sequence where it is closest to value.
- [def ep_bolfi.utility.preprocessing.OCV_from_CC_CV](#) (charge, cv, discharge, name, phases, eval_↵ points=200, spline_SOC_range=(0.01, 0.99), spline_order=2, spline_smoothing=2e-3, spline_print=None, parameters_print=False)
Tries to extract the OCV curve from CC-CV cycling data.
- [def ep_bolfi.utility.preprocessing.calculate_desired_voltage](#) (solution, t_eval, voltage_scale, overpotential, three_electrode=None, dimensionless_reference_electrode_location=0.5, parameters={})
- [def ep_bolfi.utility.preprocessing.solve_all_parameter_combinations](#) (model, t_eval, parameters, parameters_to_try, submesh_types, var_pts, spatial_methods, full_factorial=True, **kwargs)
- [def ep_bolfi.utility.preprocessing.prepare_parameter_combinations](#) (parameters, parameters_to_try, covariance, order_of_parameter_names, transform_parameters, confidence)
Calculates all permutations of the parameter boundaries.
- [def ep_bolfi.utility.preprocessing.parallel_simulator_with_setup](#) (model, current_input, parameters, submesh_types, var_pts, spatial_methods, calc_esoh, inputs, t_eval, voltage_scale, overpotential, three_↵ electrode, dimensionless_reference_electrode_location, kwargs)
- [def ep_bolfi.utility.preprocessing.simulate_all_parameter_combinations](#) (model, current_input, submesh_↵ _types, var_pts, spatial_methods, parameters, parameters_to_try=None, covariance=None, order_of_↵ parameter_names=None, additional_input_parameters=[], transform_parameters={}, confidence=0.↵ 95, full_factorial=True, calc_esoh=False, voltage_scale=1.0, overpotential=False, three_electrode=None, dimensionless_reference_electrode_location=0.5, **kwargs)

7.13 utility/visualization.py File Reference

Namespaces

- [ep_bolfi.utility.visualization](#)

Various helper and plotting functions for common data visualizations.

Functions

- def [ep_bolfi.utility.visualization.update_limits](#) (ax, xmin=float('inf'), xmax=-float('inf'), ymin=float('inf'), ymax=-float('inf'))
Convenience function for adjusting the view.
- def [ep_bolfi.utility.visualization.set_fontsize](#) (ax, title=12, xaxis=12, yaxis=12, xticks=12, yticks=12, legend=12)
Convenience function for fontsize changes.
- def [ep_bolfi.utility.visualization.update_legend](#) (ax, additional_handles=[], additional_labels=[], additional_handler_map={})
Makes sure that all items remain and all items show up.
- def [ep_bolfi.utility.visualization.push_apart_text](#) (fig, ax, text_objects, lock_xaxis=False, temp_path="./temp_render.png")
Push apart overlapping texts until no overlaps remain.
- def [ep_bolfi.utility.visualization.make_segments](#) (x, y)
Create a list of line segments from x and y coordinates.
- def [ep_bolfi.utility.visualization.colorline](#) (x, y, z=None, cmap=plt.get_cmap('viridis'), norm=matplotlib.colors.Normalize(0, 1), linewidth=1, linestyle='-', alpha=1.0)
Generates a colored line using LineCollection.
- def [ep_bolfi.utility.visualization.nyquist_plot](#) (fig, ax, ω , Z, cmap=plt.get_cmap('tab20b'), ls='-', lw=3, title_text="Impedance Measurement", legend_text="impedance", colorbar_label="Frequency / Hz", add_frequency_colorbar=True, equal_aspect=True)
Plot an impedance measurement.
- def [ep_bolfi.utility.visualization.bode_plot](#) (fig, ax_real, ax_imag, ω , Z, cmap=plt.get_cmap('tab20b'), ls_real='-', ls_imag='-', lw=3, title_text="Impedance Measurement", legend_text="impedance")
Plot an impedance measurement.
- def [ep_bolfi.utility.visualization.plot_comparison](#) (ax, solutions, errorbars, experiment, solution_visualization=[], t_eval=None, title="", xlabel="", ylabel="", feature_visualizer=lambda *args:[], feature_fontsize=12, interactive_plot=False, output_variables=None, voltage_scale=1.0, use_cycles=False, overpotential=False, three_electrode=None, dimensionless_reference_electrode_location=0.5, parameters=None)
Tool for comparing simulation<->experiment with features.
- def [ep_bolfi.utility.visualization.cc_cv_visualization](#) (fig, ax, dataset, max_number_of_clusters=4, cmap=plt.get_cmap('tab20c'), check_location=[0.1, 0.7, 0.2, 0.225])
Automatically labels and displays a CC-CV dataset.
- def [ep_bolfi.utility.visualization.plot_OCV_from_CC_CV](#) (ax_ICA_meas, ax_ICA_mean, ax_OCV_meas, ax_OCV_mean, charge, cv, discharge, name, phases, eval_points=200, spline_SOC_range=(0.01, 0.99), spline_order=2, spline_smoothing=2e-3, spline_print=None, parameters_print=False)
Visualizes the OCV_fitting.OCV_from_CC_CV output.
- def [ep_bolfi.utility.visualization.plot_ICA](#) (ax, SOC, OCV, name, spline_order=2, spline_smoothing=2e-3, sign=1)
Show the derivative of charge by voltage.
- def [ep_bolfi.utility.visualization.plot_measurement](#) (fig, ax, dataset, title, cmap=plt.get_cmap('tab20c'), plot_current=True)
Plots current and voltage curves in one diagram.

- def [ep_bolfi.utility.visualization.fit_and_plot_OCV](#) (ax, SOC, OCV, name, phases, SOC_range_↵
bounds=(0.2, 0.8), SOC_range_limits=(0.0, 1.0), z=1.0, T=298.15, fit=None, eval_SOC=[0, 1], eval_↵
points=200, spline_SOC_range=(0.01, 0.99), spline_order=2, spline_print=None, parameters_print=False,
inverted=True, info_accuracy=True, normalized_xaxis=False, distance_order=2, weights=None, initial_↵
parameters=None, minimize_options=None)

Fits an SOC(OCV)-model and an OCV(SOC)-evaluable spline.

- def [ep_bolfi.utility.visualization.visualize_correlation](#) (fig, ax, correlation, names=None, title=None,
cmap=plt.get_cmap('BrBG'), entry_color='w')

Produces a heatmap of a correlation matrix.