aiida-aimall: A Python package for automating workflows for AIMAll software

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Summary

Since its introduction by Richard Bader, the Quantum Theory of Atoms in Molecules (QTAIM) has become a useful tool for computational chemists. This Python package provides plugins for a common QTAIM software, AIMAll, for the AiiDA Python infrastructure. `aiida-aimall` is an essential tool for ensuring reproducible calculations, with full generation history. Workflows are also provided to interface `AIMAll` software with any quantum chemistry package that can be run through the command line, so long as it generates the input files required by `AIMAll`.

Statement of need

`aiida-aimall` is a Python package based on the AiiDA (Talirz et al., 2020) infrastructure designed to assist users with generating inputs for AIMAll software (Keith, 2019). The goal of the AiiDA infrastructure are, in part, to ensure data provenance and calculation reproducibility. While `aiida-aimall` has been developed primarily for interface with Gaussian software outputs (Frisch et al., 2016), through modification of classes provided by `aiida-gaussian` (Eimre et al., 2023), a versatile workflow enabling interface with other quantum chemistry packages is also made available.

Through a variety of workflows that can start with Cartesian coordinates, or even with a SMILES string of a molecule, `aiida-aimall` provides a variety of use cases for automating and complex workflows. Additionally tools to ensure that computers are not overloaded through too many simultaneous processes are made availabe through classes of `FromGroupSubmissionControllers` from `aiida-submission-controller` to limit active processes.

Features

`aiida-aimall` contains many different classes from `aiida` tailored to ensure ease of use of AIMAll calculations. Numerous features provided by `aiida-aimall` are described in full on the documentation webpage hosted on ReadTheDocs. A brief description of main features is provided here.

Running Simple AIMAll Calculations

The simplest functionality provided by `aiida-aimall` is running AIMAll calculations. All AIMAll calculations utilize the `AimqbParameters` Data type provided by `aiida-aimall`. The `AimqbParameters` datatype is a validator for AIMAll command line input. Command line parameters are to be provided as a dictionary, then `AimqbParameters` ensures that the parameters match options available for AIMAll software as defined on the software website, and that the correct data type is provided for each parameter. In this way, `AimqbParameters` verifies the provided input to AIMAll calculations prior to launch of the calculation. These parameters,

38 along with `SinglefileData` of a valid AIMAll input file, a `Code` object for AIMAll software,
39 and relevant metadata are provided to an `AimqbCalculation`.

40 This functionality in itself is an overcomplication of the simple process of running the software
41 normally. However, it does have some benefits. The output is already extracted and stored in
42 the database in a readily useable manner. Related, it is now simple to see the history of the
43 calculation.

Integrations with Computational Chemistry Software

45 `aiida-aimall`'s main draw is that it enables automation to link the outputs of standard
46 computational chemistry software directly to an AIMAll calculation. A list of provided workflows
47 is shown in Table COMPLETE. The software with the most robust implementation is Gaussian
48 software,([Frisch et al., 2016](#)) as Gaussian already has an implemented `aiida` package.

49 Table 1: Main workflows provided by `aiida-aimall`, their `aiida` entry points that can be used
50 to load them by `aiida.plugins.WorkflowFactory`, a brief description, and their main inputs.
51 These workflows all end with the output of an `AimqbCalculation` as their main output.

Workflow	Entry Point	Purpose	Input
———	:———:	:——-:	:——:
`QMToAIMWorkchain`	`aimall.qmtoaim`	Run a general computational chemistry software and link it to an AIMAll calculation	Shell Command Line Output File(s) to Get Shell Code
`GenerateWFXToAIMWorkchain`	`aimall.wfxtoaim`	Take non-standard AIMAll input files, and run AIMAll	.cp2k or .molden files
`GaussianToAIMWorkChain`	`aimall.g16toaim`	Run a Gaussian calculation and automatically run an AIMAll calculation on its outputs	`GaussianCalculation` inputs wfx filename
`SubstituentParameterWorkChain`	`aimall.subparam`	Compute substituent properties defined by the authors obtained from multistep calculation automatically	Input structure as .xyz file `SinglefileData`, `StructureData` or SMILES `Str` `GaussianCalculation` inputs

Controllers to limit computer burden when running large numbers of jobs

62 The last main contribution of `aiida-aimall` is through the definition of `FromGroupSubmissionControllers`
63 from the `aiida-submission-controller` package. These controllers limit active processes
64 and can be used together as demonstrated in (the example notebook) to automate the
65 entire `SubstituentParameterWorkchain`. These use a number of Workchains developed just
66 for their use in these controllers. The process flows as `SmilesToGaussianController` ->
67 `AIMAllReorController` -> `GaussianController` -> `AIMAllController`. The latter two con-
68 trollers can also be seen and used as general use controllers wrapping `GaussianCalculations`
69 and `AimqbCalculations`

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