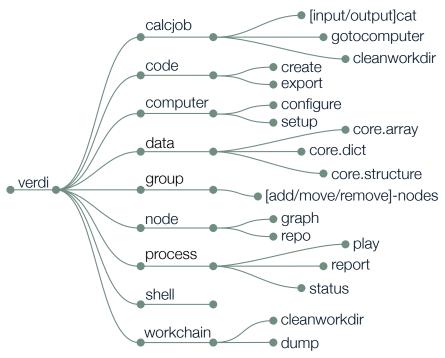
The AiDA cheat sheet

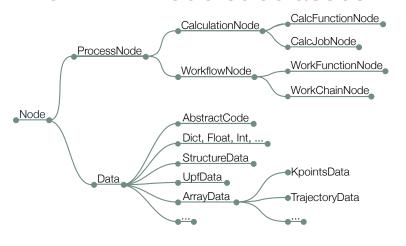


The verdi command-line API*

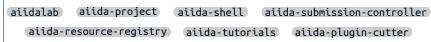


^{*}Not exhaustive

The AiiDA Node subclasses



Additional web resources (click me)



Tools of the trade

Other verdi tips and tricks

Know what's there:

- \$ verdi profile list
- \$ verdi user list
- \$ verdi storage list
- \$ verdi plugin list aiida.calculations
- \$ verdi plugin list aiida.workflows

AiiDA to classical file tree:

- \$ verdi calcjob dump <pk>
- \$ verdi workchain dump <pk>

Config options, e.g. caching:

- \$ verdi config list
- \$ verdi config set \
 - caching.default_enabled true
- \$ verdi config set caching.enabled_for \
 aiida.calculations:quantumespresso.pw

Fix what went astray:

- \$ verdi daemon restart --reset
- \$ verdi devel rabbitmq tasks analyze --fix

Share your data:

- \$ verdi archive create <.aiida-archive> \
 - --groups/--nodes <groups/nodes>
- \$ verdi archive import <.aiida-archive>

AiiDA Python imports

ORM, nodes, and Factories

Import aiida-core Node classes from aiida.orm: from aiida.orm import Dict, CalcJobNode

Import Nodes via pk, UUID, or label:
my node = load node(<identifier>)

Import Data classes via the DataFactory: (Note: Prefix AiiDA core types with core)

my_kpts = DataFactory("core.array.kpoints")

Import CalcJob classes via the CalculationFactory:
my_calcjob = CalculationFactory(

"quantumespresso.pw"

Import WorkChain classes via the WorkflowFactory.

my_workflow = WorkflowFactory(

"quantumespresso.pw.bands"







^{*}Most options also implement show/list/delete





Main attributes and methods

Node	
pk	Node ID
label	Short label
uuid	Unique ID
ctime	Creation time
mtime	Modification time
<pre>get_incoming()</pre>	Get input
<pre>get_outgoing()</pre>	Get output
inputs	All inputs generator
outputs	All outputs generator
attributes	Queryable attributes
<pre>get_attribute(k)</pre>	Attribute 'k'
extras	Queryable extras
<pre>get_extra(<k>)</k></pre>	Extra 'k'
set_extra(<k>,<v>)</v></k>	Set extra k = v
<pre>get_comments()</pre>	All comments
add_comment(<c>)</c>	Add comment with
	content <c></c>
store()	Save node in DB

Code	
load_code(<id>)</id>	Load code using pk, UUID, or label
get_builder()	Return new builder using this code

StructureData	
cell	Lattice vectors
sites	Atomic sites
kinds	Species with masses,
	symbols,
pbc	Periodic bound. cond.
	along each axis
get_formula()	Chemical formula
<pre>get_cell_volume()</pre>	Compute cell volume
convert(<fmt>)</fmt>	Convert to ASE,
	pymatgen,
set_cell(<c>)</c>	Set lattice vectors
set_ase(<a>)	Create cell from ASE
set_pymatgen()	Create cell from
	pymatgen
append_atom(Add atom of type
symbols= <symb>,</symb>	'symb' at position 'p'
position=)	

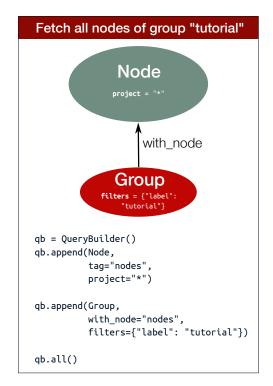
Dict	
dict. <k></k>	Get value for key 'k'
keys()	Get all keys generator
get_dict()	Get all key/values
set_dict(<dict>)</dict>	Replace all key/values

KpointsData	
<pre>set_kpoints(<k>)</k></pre>	Set an explicit list of
	kpoints 'k' (optionally
	with weights)
<pre>get_kpoints()</pre>	Get explicit list of kpts
	(if stored explicitly)
<pre>set_kpoints_mesh(</pre>	Set an implicit mesh
<m>)</m>	(e.g. 'm'=3x2x5)
<pre>get_kpoints_mesh()</pre>	Get the implicit mesh
	(if stored implicitly)

CalcJobNode	
process_state	Calc. process state
exit_status	Exit status or int code
is_finished	Has calc. finished?
is_failed	Has calc. failed?
computer	Computer where it is
	running
inputs.code	Code used to run
get_job_id()	Scheduler job ID
<pre>get_options()</pre>	Get # machines, MPI
	procs per machine,
res. <k></k>	Value of parsed
	output 'k'
	process_state exit_status is_finished is_failed computer inputs.code get_job_id() get_options()

The QueryBuilder

To import: from aiida.orm import QueryBuilder



```
Print the smearing energy calculated for BaO<sub>3</sub>Ti if it is smaller than 10<sup>-4</sup> eV
  qb = QueryBuilder()
                                                                 StructureData
  qb.append(
                                                              filters = {
    'extras.formula': 'BaO3Ti']
project = ['extras.formula']
      StructureData,
      project=["extras.formula"],
      filters={"extras.formula":"BaO3Ti"},
      tag="structure")
                                                                          with_incoming
  qb.append(
      CalcJobNode,
      tag="calculation",
                                                                 CalcJobNode
      with_incoming="structure")
  qb.append(
                                                                          with_incoming
      Dict,
      tag="results",
      filters={"attributes.energy_smearing":
                                                                      Dict
      {"<=":-0.0001}},
      project=[
           "attributes.energy_smearing",
           "attributes.energy_smearing_units"
      with_incoming="calculation")
  qb.all()
```



