

Marking up lattice QCD configurations and ensembles

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QCDml is an XML-based markup language designed for sharing QCD configurations and ensembles world-wide via the International Lattice Data Grid (ILDG). Based on the latest release, we present key ingredients of the QCDml in order to provide some starting points for colleagues in this community to markup valuable configurations and submit them to the ILDG.

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1. Introduction

Sharing QCD configurations world-wide via the ILDG [1, 2, 3, 4] requires standards of notation and terminology for metadata of configurations such as lattice actions and sizes used in simulations. For this purpose, the Metadata Working Group (MDWG) [5] has been working [2, 6, 7] on designing an XML-based markup language QCDml [8], a set of rules for XML instance documents (IDs). IDs are stored in regional grid databases which are aggregated together to form the ILDG Metadata Catalogue (MDC). Because this service is queried by users, contributors (and hopefully users) are advised to understand QCDml.

After the first working version [6] was released in 2004, the MDWG has revised the QCDml schema several times [7]. Although our design strategy and the global structure of QCDml remain unchanged, many improvements have been made to meet community requirements, some of them being incompatible with previous versions. In this article, we present key ingredients of the QCDml, based on the latest release.

2. QCDml Structure

QCDml defines the *ensemble XML* and the *configuration XML* whose structure is shown below. The former describes metadata common across an ensemble such as <physics> information, while the latter contains configuration specific information such as a trajectory number <update>, a tag <series> which distinguishes different runs in one ensemble, <precision> of the configuration, and an <avePlaquette> value. The <algorithm> parameters can be stored in either of the XML documents. The <management> part includes information of who and when data and metadata are submitted or modified. The <implementation> part contains information of machine and code used in simulations.

The two XML IDs and the configuration itself are linked together in the following way. The Markov Chain URI (MCU) in <markovChainURI> is a unique ensemble name and appears in both XML documents. The Logical File Name (LFN) in <dataLFN> of the configuration XML is a unique configuration name and is embedded in the configuration file [9]. The format of the MCU (LFN) is “mc:(lfn)/(Name of Regional Grid)/(Regional Grid Dependent String)”.

Root elements, <markovChain> for the ensemble XML and <gaugeConfiguration> for the configuration XML, contain names of namespaces (xmlns=) and URL's of schema (schemaLocation). A two digit version number of the QCDml (currently 1.4 for ensemble and 1.3 for configuration) is appended at the end of the namespace, and more one digit release number is added to the schemata file to identify backward compatible updates.

ensemble XML

```
<markovChain xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
  xsi:schemaLocation="http://www.lqcd.org/ildg/QCDml/ensemble1.4
    http://www.lqcd.org/ildg/QCDml/ensemble1.4/QCDmlEnsemble1.4.1.xsd"
  xmlns="http://www.lqcd.org/ildg/QCDml/ensemble1.4">
  <markovChainURI>
    mc://JLDG/CP-PACS+JLQCD/RCNF2+1/RC28x56_B2050Kud013560
  </markovChainURI>
```

```

<management/>
<physics/>
<algorithm/>
</markovChain>

```

configuration XML

```

<gaugeConfiguration xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
  xsi:schemaLocation="http://www.lqcd.org/ildg/QCDml/config1.3
    http://www.lqcd.org/ildg/QCDml/config1.3/QCDmlConfig1.3.0.xsd"
  xmlns="http://www.lqcd.org/ildg/QCDml/config1.3">
  <management/>
  <implementation/>
  <algorithm/>
  <precision>double</precision>
  <markovStep>
    <markovChainURI>
      mc://JLDG/CP-PACS+JLQCD/RCNF2+1/RC28x56_B2050Kud013560
    </markovChainURI>
    <series>2</series> <update>002200</update>
    <avePlaquette>6.04326596981212e-01</avePlaquette>
    <dataLFN>
      lfn://JLDG/CP-PACS+JLQCD/RCNF2+1/RC28x56_B2050Kud013560-2-002200
    </dataLFN>
  </markovStep>
</gaugeConfiguration>

```

3. Physics Part

3.1 Structure

The <physics> part of the ensemble XML consists of <size>, <action> and optional <observables>. The <size> section looks like <size> <elem> <name>X</name> <length>16</length> </elem> (lengths in Y,Z,T directions) </size>.

The <action> section is a central part of the ensemble XML. The example below illustrates the structure:

```

<action>
  <gluon> <iwasakiRGGluonAction/> </gluon>
  <quark> <npCloverQuarkAction/> <npCloverQuarkAction/> </quark>
</action>

```

Each lattice action is described in an element block (e.g. <iwasakiRGGluonAction/>) whose name is easily understood. This strategy is taken, firstly to assign an unique name to each action, and secondly to realize a hierarchal structure of action trees in the schemata. Note that we repeat quark action blocks if coupling parameters are different, e.g. for the $N_f = 2 + 1$ case. Fig. 1 shows a typical quark action. An action block consists of

- <glossary>, the URL of a document prepared by contributors using either text, pdf, ps, tex, or xml formats. The document describes full details of the action.

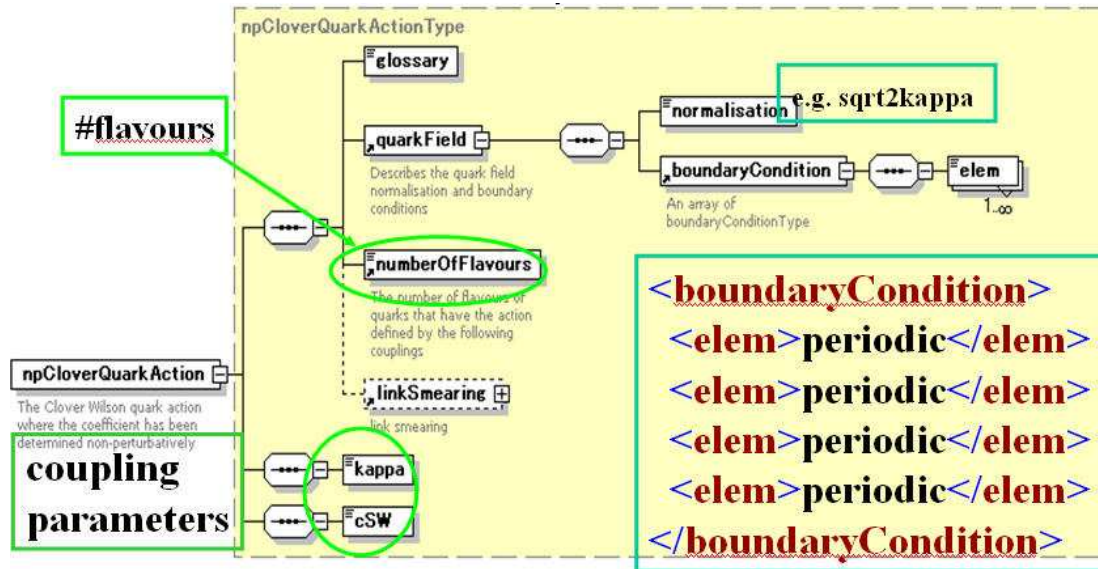


Figure 1: A graphical representation of the XML schemata for the clover quark action.

- gluon/quark field information, `<gaugeField>` or `<quarkField>`, which includes `<boundaryCondition>` and supplemental information such as `<representation>` and `<normalisation>`.
- `<numberOfFlavours>` for quark actions.
- optional `<linkSmearing>` block which is explained below.
- list of coupling parameters together with tadpole factors for tadpole improved actions.

The last optional block `<observables>` of the `<physics>` part displays values of measured physical quantities. Each element of the list has the following structure: `<elem> <name/> <value/> <err/> <glossary/> </elem>`. Currently, `<name>` of `<observables>` is one of `ampi` (am_π), `amrho` (am_ρ), `mpi_mrho` (m_π/m_ρ), `ar0` (ar_0) and `ar1` (ar_1). QCDml supports only dimensionless quantities, because dimensional quantities (e.g. lattice spacing) cannot be regarded as properties of an ensemble.

3.2 Details of actions

The current version of QCDml supports the following lattice actions (lower camel case convention).

- gluon actions: `plaquetteGluonAction`, `iwasakiRGGluonAction`, `DBW2GluonAction`, `LuscherWeiszGluonAction`, `treeLevelSymanzikGluonAction`, `tpLuscherWeiszGluonAction`, `anisotropicWilsonGluonAction`, `anisotropicTpWilsonGluonAction`
- quark actions: `KSquarkAction`, `asqTadQuarkAction`, `wilsonQuarkAction`, `cloverQuarkAction`, `tpCloverQuarkAction`, `npCloverQuarkAction`, `wilsonTMQuarkAction`, `domainWallQuarkAction`, `fatLinkIrrelevantCloverQuarkAction`, `anisotropicWilsonQuarkAction`, `anisotropicCloverQuarkAction`

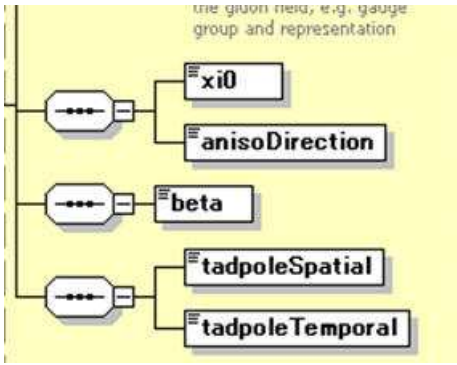


Figure 2: Coupling parameters of the anisotropic tadpole improved Wilson gluon action. $\langle xi0 \rangle$ is the bare gauge anisotropy and $\langle anisoDirection \rangle$ is the direction of anisotropy.

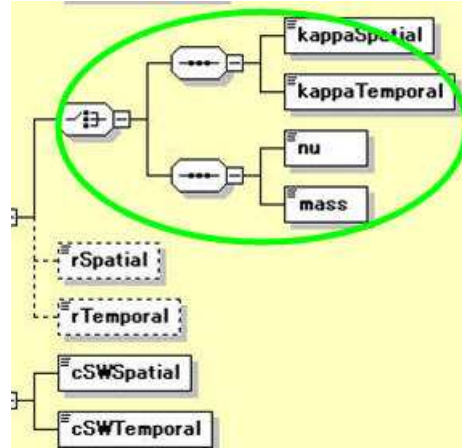


Figure 3: Coupling parameters of the anisotropic clover quark action. Wilson parameters are optional.

A list of coupling names for each action can be found in Ref. [10].

There are currently two examples of non-unique notation in QCDml.

- Gluon actions with plaquette and six-link loops (sixLinkGluonAction), such as $\langle DBW2GluonAction \rangle$, are mathematically written as

$$S = \beta \times (c_0 \sum \text{plaq.} + c_1 \sum \text{rect.} + c_2 \sum \text{parallelogram} + c_3 \sum \text{char}).$$

The QCDml supports two standard normalisation of couplings;

$$c_0 = 1, \quad c_0 + 8c_1 + 8c_2 + 16c_3 = 1.$$

The normalisation is marked up as a value of the $\langle normalisation \rangle$ element, $c0_is_one$ for the former (recommended for tadpole improved actions) and $cs_sum_to_one$ for the latter (recommended for others).

- The current version of QCDml supports some anisotropic actions. See figs. 2 and 3 for examples. Note that one can choose either $(\kappa_{\text{Spatial}}, \kappa_{\text{Temporal}})$ or (μ, mass) notations for anisotropic wilson/clover quark actions.

3.3 Link smearing

Smearing link variables is becoming one of the standard technologies for improvement. QCDml supports marking up smearing as an optional $\langle linkSmearing \rangle$ block in all quark actions. Link smearing consists of a blocking of link variables using e.g. an APE type blocking

$$U_\mu(n) \leftarrow \rho_0 U_\mu(n) + \rho_1 \sum_{v(\neq\mu)} [U_v(n)U_\mu(n+\hat{v})U_v^\dagger(n+\hat{\mu}) + U_v^\dagger(n-\hat{v})U_\mu(n-\hat{v})U_v(n-\hat{v}+\hat{\mu})]$$

and a projection onto the SU(3) group (unitarization) using e.g. a stout procedure. One repeats the set of “blocking” and “unitarization” several times. QCDml therefore marks up link smearing procedures as

```

<linkSmearing>
  <apeLinkBlocking> <rho0/> <rho1/> </apeLinkBlocking>
  <stoutLinkUnitarization/>
  <numSmear/>
</linkSmearing>

```

In addition to the above example, we currently support `<anisotropicApeLinkBlocking>` and `<invSqrtLinkUnitarization>`. The strategy of inserting an optional `<linkSmearing>` block to all actions makes it easy to markup various fat link actions.

4. Algorithm Part

Marking up algorithms is realized with a strategy different from that for actions. Due to the variety and complexity of algorithms, a generic hierarchical markup is difficult. Moreover, it is unlikely that the algorithm would form the primary search criteria. Therefore, algorithms are marked up in a rather unconstrained way.

The algorithm part in the ensemble XML looks like `<algorithm> <name/> <glossary/> <reference/> <exact/> <parameters/> </algorithm>`. The name of the algorithm is given by contributors as a value of the `<name>` element. `<glossary>` and `<reference>` elements, also unconstrained, refer to the URL of a document and an article reference, respectively. The only constrained element `<exact>` provides information of whether the algorithm is exact (true) or not (false). The `<parameters>` block contains a list of `<name>` and `<value>` pairs like

```

<parameters>
  <name>PHMC_Polynomial_Order</name>    <value>250</value>
  <name>PHMC_Polynomial_Type</name>     <value>Chebyshev</value>
  <name>HMC_MD_dt</name>                 <value>0.400000000E-02</value>
  <name>HMC_MD_steps</name>             <value>250</value>
</parameters>

```

Algorithm parameters might be changed when generating an ensemble. For example, the molecular dynamics step size of HMC is often tuned in a run. Contributors can mark up such configuration dependent parameters in the `<algorithm>` part of the configuration XML.

In addition to the `<parameters>` list, contributors can import a collaboration specific namespace and mark up algorithm parameters in a hierarchical way. Information given in this way is utilized only within the collaboration.

5. Management Part

The `<management>` part of the ensemble XML describes information for the XML document itself. This part includes `<collaboration>`, `<projectName>`, optional `<ensembleLabel>` and `<reference>`. These elements provide users with a starting point to search useful ensembles. The `<management>` part of the configuration XML includes `<crcChecksum>` which is a crc check sum of the binary data part of the configuration file [9]. Checking this value ensures that the configuration is successfully transferred from the ILDG.

In addition, we can record operations (add, replace, remove) affecting an ensemble XML document in the MDC together with the time stamp and the participant responsible for the operation.

Similarly, operations (generate, add, replace, remove) affecting a configuration (or configuration XML metadata) are recorded in the <management> part of the configuration XML. Marking up these operations is optional, except for the information when a configuration has been generated.

6. Summary and Future Work

At present, interoperability of the regional grids has been achieved [11] for download operations and valuable configurations have already been archived [4] in the grid. Because the ILDG is becoming a new research infrastructure for the lattice QCD community, the MDWG encourages our colleagues to mark up their ensembles and configurations using QCDml.

The description of QCDml given in this report is not complete. To study more about QCDml, please visit the ILDG MDWG web site [10] where you can find sample XML documents together with human readable documents generated from annotations embedded in the QCDml schema.

The lattice actions used in simulations are becoming more and more complicated. We plan to prepare human readable documents including mathematical expressions and reference articles in the near future. We are going to update QCDml continuously to meet community requirements.

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