#

# **README**

Title: README.md (examples-emdb) Author: jdw Date: 17-Feb-2017 Updated: 19-Feb-2017

## **Example Files**

Example files for the EMDB content request service.

## **Example Service Content Type Definitions**

#### **Example Summary Content Definition**

The following JSON snippet describes an example summary content type definition. The content items refer to table and column names in the various OneDep database schema.

```
"author list emdb",
        "exp_method",
        "status_code_exp",
        "emdb_id",
        "status_code_emdb",
        "dep_author_release_status_code_emdb"
    "em_admin": [
        "entry_id",
        "structure id",
        "title",
        "map release date",
        "deposition date"
},
"resource": {
    "deposition": [
        "status",
        "status"
    "em_admin": [
        "da_internal",
        "da internal"
},
"conditions": {
    "deposition": {
        "exp_method": [
            "ELECTRON MICROSCOPY",
            "char",
            "LIKE"
},
"type": "rdbms"
```

### **Example Entry-level Content Definition**

2/19/2017 README

The following JSON snippet describes an example entry-level content type definition. The content items refer to PDBx/mmCIF category and attribute names. This content definition may be applied to data files within both the deposition and biocuration systems, dialects for both EM namespaces are included in this example.

```
"report-entry-example-emdb": {
      "content": {
          "pdbx database status": [
              "status code",
              "author_release_status_code",
              "deposit site",
              "process site",
              "recvd initial_deposition_date",
              "date of NDB release"
          "entity poly": [
              "entity id",
              "type",
              "nstd linkage",
              "nstd monomer",
              "pdbx seq one letter code",
              "pdbx seq one letter code can"
          "emd_admin": [
              "entry_id",
              "current status",
              "deposition date",
              "deposition site",
              "details",
              "map hold date",
              "last update",
              "map release date",
              "obsoleted date",
              "replace existing entry flag",
              "title".
              "header release date"
          "database 2": [
              "database id",
              "database code"
```

README

2/19/2017

```
"em map": [
    "id",
    "annotation details",
    "axis_order_fast",
    "axis_order_medium",
    "axis_order_slow",
    "cell_a",
    "cell_alpha",
    "cell_b",
    "cell_beta",
    "cell_c",
    "cell gamma",
    "contour level",
    "contour level source",
    "data type",
    "dimensions_col",
    "dimensions_row",
    "dimensions_sec",
    "endian_type",
    "file",
    "format",
    "label",
    "limit col",
    "limit row",
    "limit_sec",
    "origin_col",
    "origin row",
    "origin_sec",
    "partition",
    "pixel_spacing_x",
    "pixel_spacing_y",
    "pixel spacing z",
    "size_kb",
    "spacing x",
    "spacing_y",
    "spacing_z",
    "statistics_average",
    "statistics maximum",
```

README

```
"statistics_minimum",
    "statistics_std",
    "symmetry_space_group",
    "type"
"entity": [
   "id",
    "type",
    "src method",
    "pdbx_description",
    "pdbx number of molecules"
"struct": [
    "entry_id",
    "title"
"em_admin": [
    "entry_id",
    "current_status",
    "deposition_date",
    "deposition_site",
    "details",
    "map_hold_date",
    "last update",
    "map_release_date",
    "obsoleted date",
    "replace_existing_entry_flag",
    "title",
    "header release_date"
"emd_map": [
    "id",
    "contour level",
    "contour level source",
    "annotation details",
    "file",
    "original file",
    "label",
    "type",
```

```
"partition",
    "format",
    "size kb",
    "axis_order_fast",
    "axis_order_medium",
    "axis_order_slow",
    "cell alpha",
    "cell_beta",
    "cell gamma",
    "cell_a",
    "cell b",
    "cell_c",
    "data_type",
    "dimensions_col",
    "dimensions_row",
    "dimensions_sec",
    "origin_col",
    "origin_row",
    "origin_sec",
    "limit col",
    "limit row",
    "limit sec",
    "pixel_spacing_x",
    "pixel_spacing_y",
    "pixel_spacing_z",
    "symmetry_space_group",
    "spacing x",
    "spacing_y",
    "spacing z",
    "statistics_minimum",
    "statistics maximum",
    "statistics_average",
    "statistics std",
    "endian type"
"audit author": [
    "name",
    "pdbx ordinal"
```