BIGSdb Isolate Database (version 1.27) projects Isolates Loci PK user_name composite field values composite fields PK.FK1 short description record action PK,FK2 composite field order PK,FK1 PK,FK1 PK,FK1 <u>isolate id</u> PK,FK2 <u>field</u> isolate id field isolate display modify data_type main_display private empty_value locus_type allele_id_format curator curator no_quota regex curator primer1 primer2 restrict_user restrict_usergro Sequences pcr_locus FK1 allele_id_regex and max_length eav_fields common name FK1 project members curator base uri designations PK description url isolate_value_extended_attributes FK3 curator PK,FK2 isolate id length_varies coding_sequence remote contias PK,FK1 access token access secret datestamp PK.FK1 isolate field session secret PK,FK1 isolate id probes FK1 value_regex conditional_formatting pcr_filter probe_filter match_longest dbase_name length checksum date_entered probe locus FK2 curator html_link_text project user groups PK,FK1 PK,FK2 html message dbase_host PK,FK1 project id user group max_value field_order dbase_port isolates description curator FK1 dbase password no_curate min_alignment max_mismatch PK isolate field extended attributes PK isolate id max gaps datestamn main_display isolate_display query_field analysis isolate_id curator (FK1 FK3 locus aliases country FK1 curator experiment_sequences year age sex disease PK,FK1 value_format eav text experiment ic isolate aliases run id value regex PK,FK2 PK,FK1 <u>isolate id</u> PK,FK2 <u>field</u> original_designation isolate id FK1 source curator curator date entered length field_order FK2 serogroup comments value FK3 FK2 curator curator PK,FK1 isolate id new_version sender 丰 FK1 FK2 curator accession history PK,FK1 sequence attribute values Classification groups eav date PK.FK1 databank id PK.FK1 PK.FK2 isolate id PK,FK1 isolate id PK.FK2 kev FK2 classification_schemes PK.FK2 curator action curator bookmarks FK2 value validation_conditions PK classification group fields validation_rule_conditions FK2 scheme id PK,FK2 PK validation_rules PK,FK1 PK,FK2 PK,FK1 id flag dbase config description condition is set_id inclusion_threshold PK operator value sequence_attributes page params seqdef_cscheme_id description field_order datestam FK3 FK2 PK curator FK1 FK1 user id curator FK1 type descriptio FK1 last accessed datestamp public allele_sequences FK1 allele_designations curator PK ⊕ FK5 isolate_id FK4 isolate id Schemes 0 FK1 FK2 Users seqbin_id set loci FK1 scheme members locus start_pos scheme_flags FK2 seabin stats scheme warehouse indices Ŧ status OF FK4 PK,FK2 PK,FK2 isolate id PK,FK1 PK,FK1 set_name formatted_set_name FK3 PK,FK2 field_order date_entered FK2 O€ FK3 FK3 PK datestamp formatted set common name display orde user name dbase_name dbase_host first name 考 schemes Sets dbase_port submissions **Submissions** PK.FK1 PK,FK1 dbase user dbase_password scheme_group_group_members PK.FK3 PK list_order user id account request emails description dbase name type primary_key description url auto_registration FK1 link order PK.FK1 isolate_submission_field_orde PK,FK2 FK1 description FK2 curator datestamp metadata id datestamp long_descripti date_entered PK,FK1 PK FK2 dbase_port dbase_user display_order hidden FK2 curator field order main_display isolate_displa query_field dropdown FK1 FK2 dbase_password curator curator index user_groups dbase id outcome scheme_refs user group members PK <u>id</u> isolate display permissions FK2 curator PK,FK1 PK,FK2 PK,FK1 query_status description PK,FK1 scheme group scheme members set view messages PK.FK2 set id analysis co_curate_private -**()**← FK3 PK,FK1 PK,FK2 PK,FK1 PK,FK1 submission id PK,FK1 set_id display_order curator index field allow_missing_loci FK1 curator FK1 curator FK3 curator FK2 curator FK2 datestamp Curator field foreign key relationships to users table are not shown to aid clarity