

schema

| hpfinding | organweights | ccfinding | clinicalhaematologicalfindings |
|--|-------------------|-------------------------------|---|
| subst_id | subst_id | subst_id | subst_id |
| study_id | study_id | study_id | study_id |
| dose | dosemgkg | dose | dosemgkg |
| sex | sex | sex | sex |
| standardisedsex | standardisedsex | standardisedsex | standardisedsex |
| normalised_sex | normalised_sex | normalised_sex | normalised_sex |
| relevance | relevance | relevance | relevance |
| grade | | | |
| | sd | sd | sd |
| | averagefoldchange | average_value | averagefoldchange |
| | averageweight | average_fold_change | averagevalue |
| | unit | unit | unit |
| timepoint | timepoint | timepoint | timepoint |
| timepoint_unit | timepointunit | timepoint_unit | timepointunit |
| | finding | finding | finding |
| | | | |
| histopathology_organ_affected | organweighted | | |
| histopathology_organ_affected_standardised | standardisedorgan | | |
| histopathology_organ_affected_standardised_revised | normalised_organ | | |
| histopathology_finding | | clinical_chemistry_parameter | clinicalhaematologyparameter |
| histopathology_finding_standardised | | standarised_parameter | standardisedparameter |
| histopathology_finding_standardised_revised | | standarised_parameter_revised | normalised_clinicalhaematologyparameter |
| | | | |
| num_animals_total | | | |
| num_animals_affected | | | |
| | | | |
| eliminar source | | eliminar source | |

schema

urinalysisfindings

subst_id
study_id
dosemgkg
sex
standardisedsex
normalised_sex
relevance

sd
averagefoldchange
averagevalue
unit
timepoint
timepointunit

finding

urinalysisparameter
standardisedparameter
normalised_urinalysisparameter

subst_id
study_id
dose
sex
standardisedsex
normalised_sex
relevance
grade
sd
averagefoldchange
averagevalue
unit
timepoint
timepointunit

change

organ_verbatim
organ_standardised
organ_normalised
finding_verbatim
finding_standardised
finding_normalised

num_animals_total
num_animals_affected

source