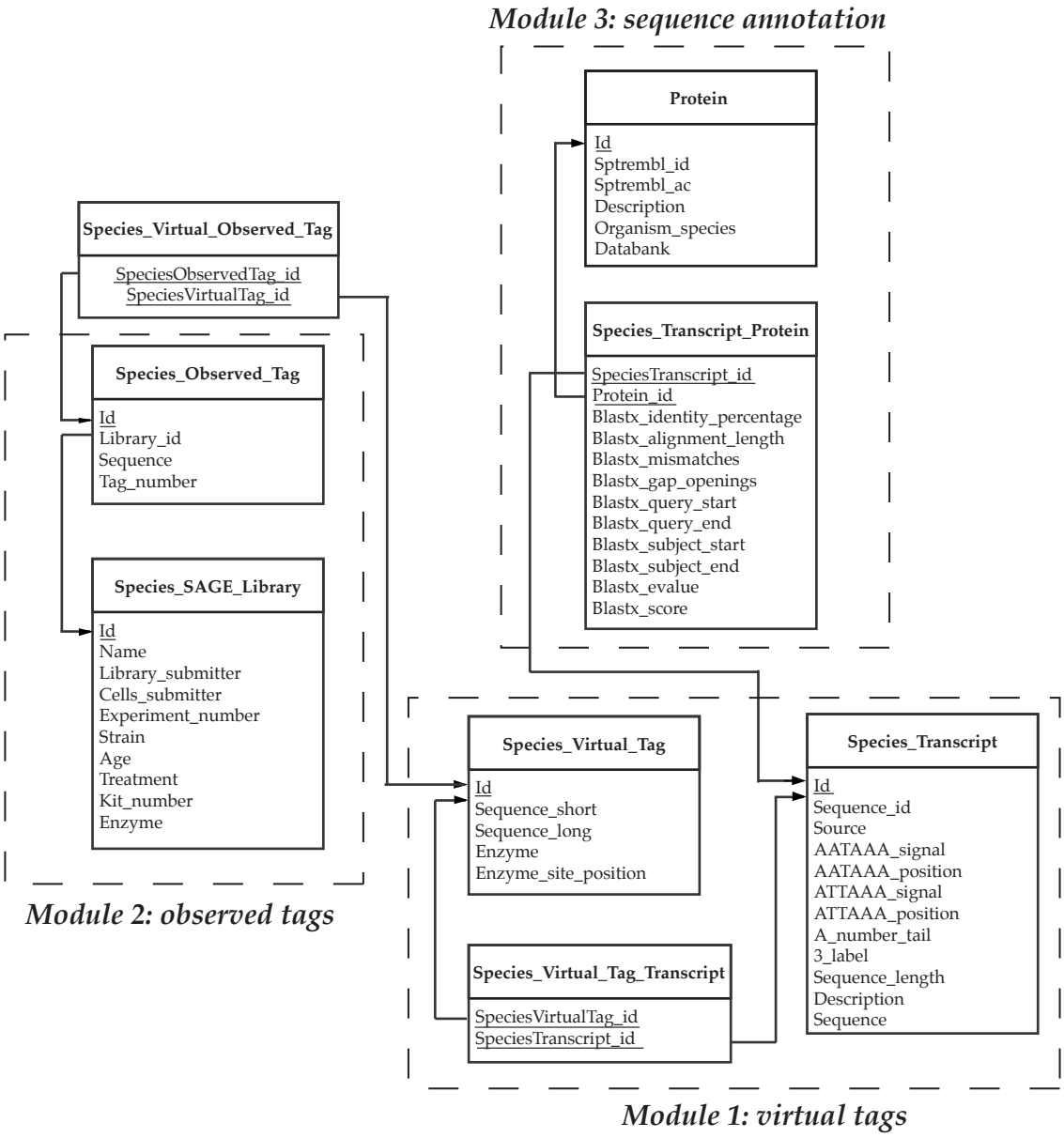
Module 3: sequence annotation
Protein
Id
Sptrembl_id
Sptrembl_ac
Description
Organism_species
Databank
Species_Transcript_Protein
SpeciesTranscript_id
Protein_id
Blastx_identity_percentage
Blastx_alignment_length
Blastx_mismatches
Blastx_gap_openings
Blastx_query_start
Blastx_query_end
Blastx_subject_start
Blastx_subject_end
Blastx_evalue
Blastx_score
Species_Virtual_Observed_Tag
SpeciesObservedTag_id
SpeciesVirtualTag_id
Species_Observed_Tag
Id
Library_id
Sequence
Tag_number
Species_SAGE_Library
Id
Name
Library_submitter
Cells_submitter
Experiment_number
Strain
Age
Treatment
Kit_number
Enzyme
Module 2: observed tags
Species_Virtual_Tag
Id
Sequence_short
Sequence_long
Enzyme
Enzyme_site_position
Species_Virtual_Tag_Transcript
SpeciesVirtualTag_id
SpeciesTranscript_id
Species_Transcript
Id
Sequence_id
Source
AATAAA_signal
AATAAA_position
ATTAAA_signal
ATTAAA_position
A_number_tail
3_label
Sequence_length
Description
Sequence
Module 1: virtual tags