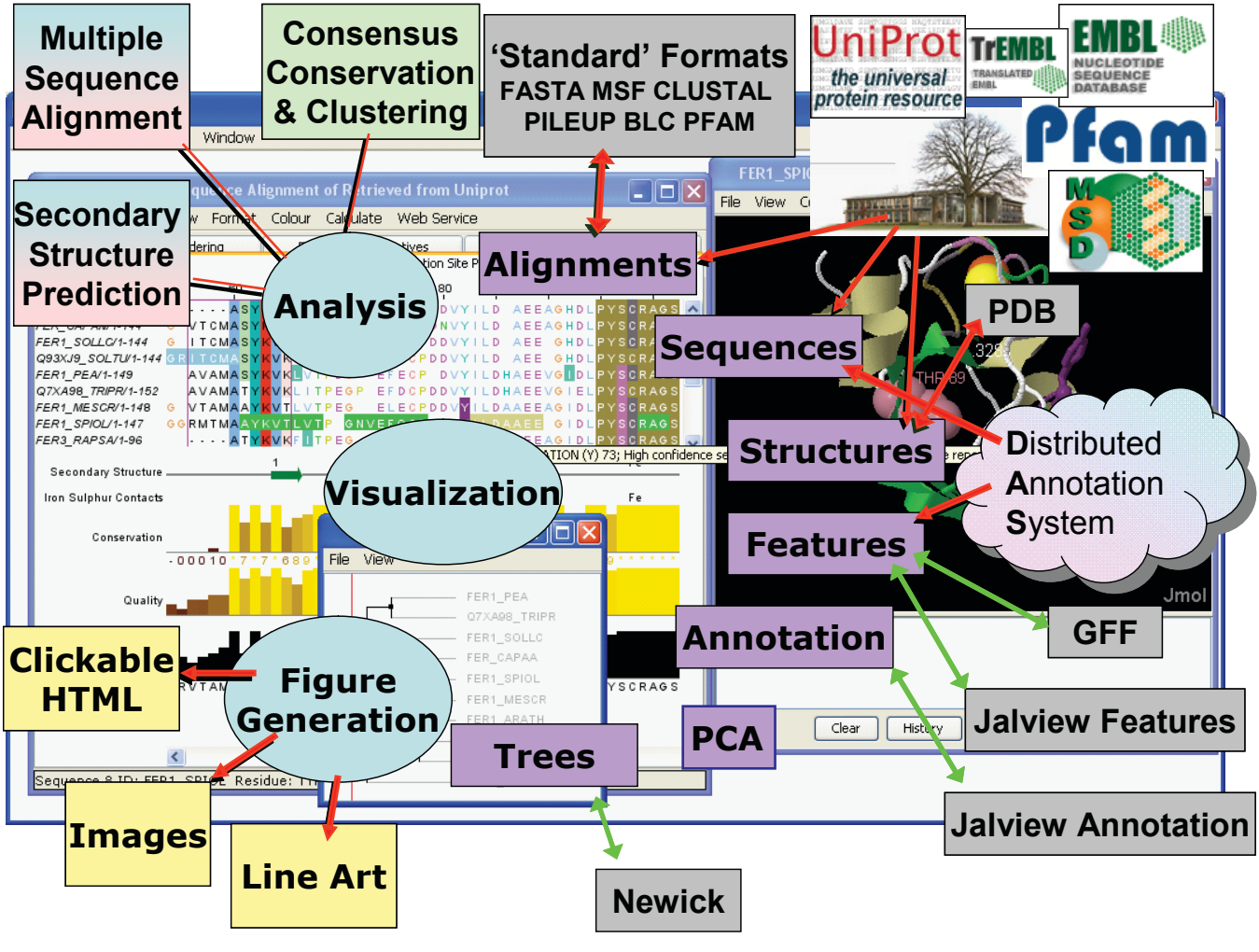
Multiple Sequence Alignment
Consensus Conservation & Clustering
'Standard' Formats
FASTA MSF CLUSTAL
PILEUP BLC PFAM
UniProt
the universal protein resource
TrEMBL
EMBL
NUCLEOTIDE SEQUENCE DATABASE
Pfam
MSD
Secondary Structure Prediction
Analysis
Alignments
Sequences
PDB
Structures
Distributed Annotation System
Features
Visualization
Annotation
GFF
Clickable HTML
Figure Generation
PCA
Jalview Features
Trees
Jalview Annotation
Images
Line Art
Newick
Window
Format
Colour
Calculate
Web Service
Secondary Structure
Iron Sulphur Contacts
Conservation
Quality
File
View
Clear
History
Jmol