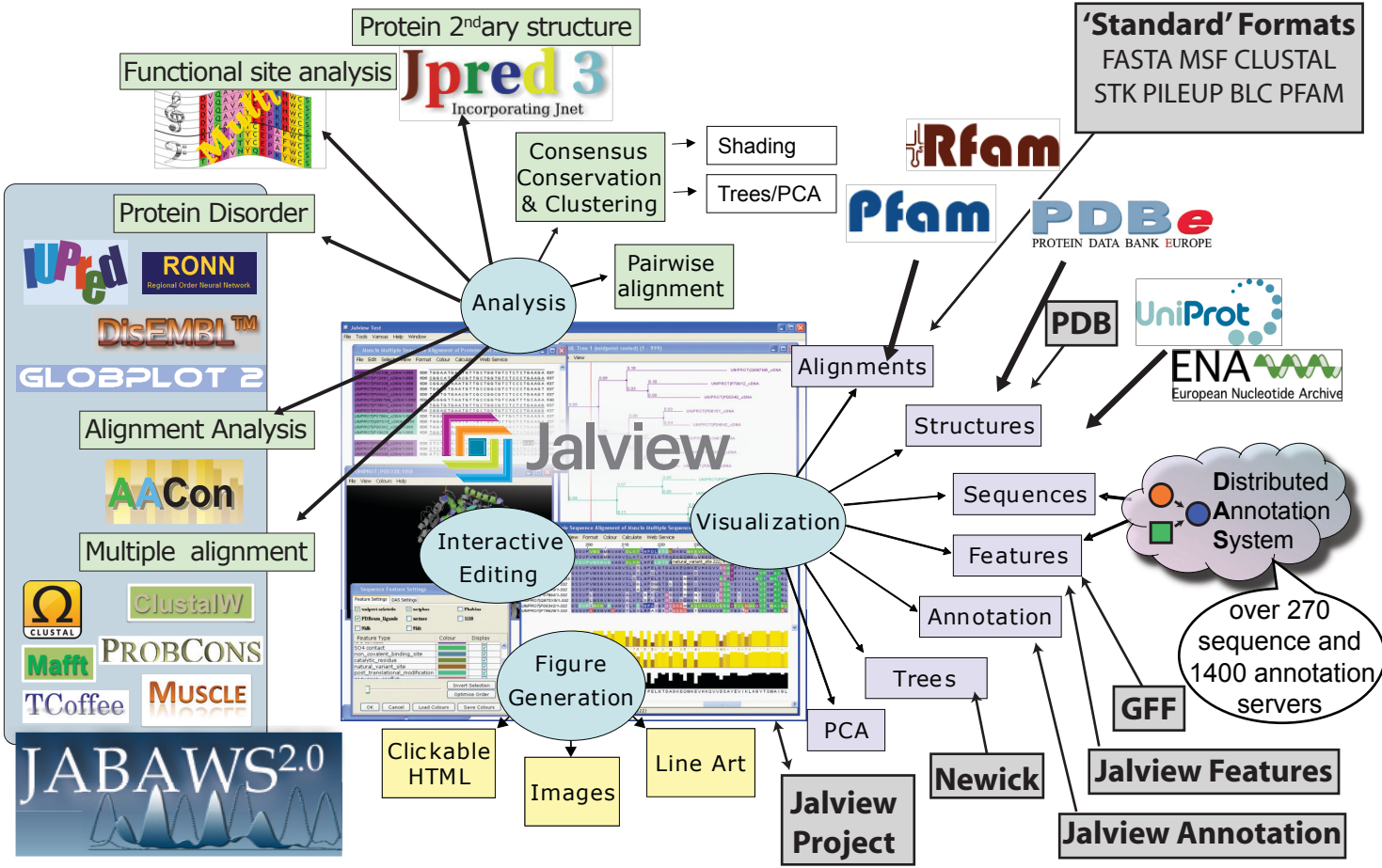
Protein 2ndary structure
Jpred 3
Incorporating Jnet
Functional site analysis
Consensus Conservation & Clustering
Shading
Trees/PCA
'Standard' Formats
FASTA MSF CLUSTAL
STK PILEUP BLC PFAM
Rfam
Pfam
PDBe
PROTEIN DATA BANK EUROPE
Protein Disorder
IUPred
RONN
Regional Order Neural Network
DisEMBL™
GLOBPLOT 2
Analysis
Pairwise alignment
PDB
UniProt
ENA
European Nucleotide Archive
Alignments
Structures
Alignment Analysis
AACon
Jalview
Visualization
Sequences
Distributed Annotation System
Features
Interactive Editing
Multiple alignment
Annotation
ClustalW
CLUSTAL
Mafft
PROBCONS
TCoffee
MUSCLE
Figure Generation
Trees
over 270 sequence and 1400 annotation servers
GFF
PCA
Clickable HTML
Images
Line Art
Newick
Jalview Features
Jalview Project
Jalview Annotation
JABAWS2.0