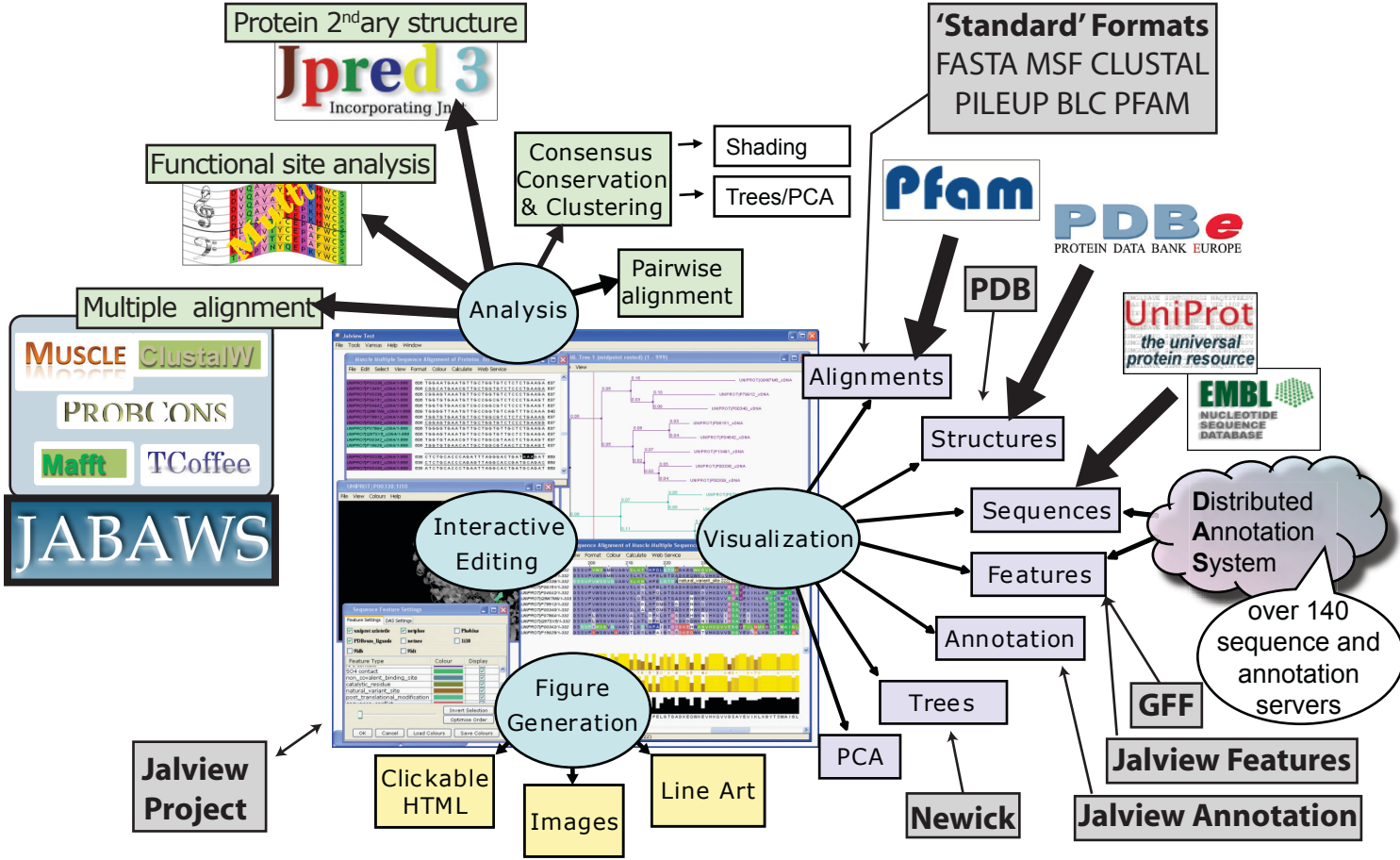
Protein 2ndary structure
Jpred 3
Incorporating Jn
Functional site analysis
Consensus Conservation & Clustering
Shading
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Multiple alignment
MUSCLE
ClustalW
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Analysis
'Standard' Formats
FASTA MSF CLUSTAL
PILEUP BLC PFAM
Pfam
PDBe
PROTEIN DATA BANK EUROPE
PDB
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the universal protein resource
EMBL
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over 140 sequence and annotation servers
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