

PDB Batch v1 – Results Explained

PDB Batch v1 explains its internal logic through the results it produces. The sequence comparison reveals whether structural changes exist between wild-type and mutant proteins. Mutation tables enumerate substitutions, insertions, and deletions with positional accuracy. The color-mapped sequence view translates alignment logic into an intuitive visual language, while physicochemical profiles explain how mutations alter protein behavior. Ramachandran plots validate structural integrity, and 3D visualization confirms spatial consequences of mutations.