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| **Data provider** | **Annotation Type** | **Reference** |
| 14-3-3-pred | Predicted 14-3-3-binding sites | ([Madeira et al., 2015](#_ENREF_10)) |
| 3D-Complex | Interaction interface annotations | ([Levy, Pereira-Leal, Chothia, & Teichmann, 2006](#_ENREF_9)) |
| 3D-LigandSite  | Putative ligand binding sites | ([Wass, Kelley, & Sternberg, 2010](#_ENREF_20)) |
| AKID | Predicted kinase-specific phosphorylation | ([Parca et al., 2019](#_ENREF_12)) |
| CaMKinet | Calmodulin-dependent kinase annotations | (*in preparation*) |
| canSAR | Predicted druggable pockets | ([Coker et al., 2019](#_ENREF_3)) |
| CATH-FunSites | Predicted functional sites | ([Ashford, Pang, Moya-Garcia, Adeyelu, & Orengo, 2019](#_ENREF_1)) |
| ChannelsDB  | Predicted molecular channels | ([Pravda et al., 2018](#_ENREF_13)) |
| Covalentizer | Potential covalent binder sites | (*in preparation*) |
| DEPTH | Residue depth | ([Tan, Nguyen, Patel, Varadarajan, & Madhusudhan, 2013](#_ENREF_18)) |
| DynaMine | Predicted backbone flexibility | ([Cilia, Pancsa, Tompa, Lenaerts, & Vranken, 2013](#_ENREF_2)) |
| EFoldMine | Predicted early folding residues | ([Raimondi, Orlando, Pancsa, Khan, & Vranken, 2017](#_ENREF_15)) |
| FireProt DB | Effect on the stability of point mutations | ([Stourac et al., 2021](#_ENREF_17)) |
| FoldX | Effect on the stability of point mutations | ([Delgado, Radusky, Cianferoni, & Serrano, 2019](#_ENREF_5)) |
| KinCore | Conformational annotations of kinases | (*in preparation*) |
| KnotProt | Topology annotations | ([Dabrowski-Tumanski et al., 2019](#_ENREF_4)) |
| M-CSA  | Catalytic sites | ([Ribeiro et al., 2018](#_ENREF_16)) |
| MetalPDB | The biological relevance of metal ions | ([Putignano, Rosato, Banci, & Andreini, 2018](#_ENREF_14)) |
| Missense3D | Effect on the stability of point mutations | ([Ittisoponpisan et al., 2019](#_ENREF_6)) |
| P2rank  | Predicted ligand binding sites | ([Krivak & Hoksza, 2018](#_ENREF_8)) |
| POPScomp | Accessible surface area | ([Kleinjung & Fraternali, 2005](#_ENREF_7)) |
| ProKinO | Protein kinase annotations | ([McSkimming et al., 2015](#_ENREF_11)) |
| WEBnm@ | Flexibility predictions | ([Tiwari et al., 2014](#_ENREF_19)) |

**Supplementary Table 1 - PDBe-KB partner resources providing annotations**

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