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| **S6 Table. Structural templates used by ITASSER to model AvrE and HaRxL23 structures** | | |
| **AvrE templates** | **HaRxL23 templates** |
| 4IGL: Structure of the RHS-repeat containing BC component of the secreted ABC toxin complex from Yersinia entomophaga | 5L7S2: Crystal structure of RXLR effector PexRD54 from Phytophthora infestans |
| 5A9Q: Human nuclear pore complex | 2LC2: Solution structure of the RXLR effector P. capsici AVR3a4 |
| 5YFP: Cryo-EM Structure of the Exocyst Complex | 2RGH: Structure of Alpha-Glycerophosphate Oxidase from Streptococcus sp.: A Template for the Mitochondrial Alpha-Glycerophosphate Dehydrogenase |