

Table 1. The number of protein sequences obtained from each procedure.

Source	Sequence number
Aragunde et al. (2018)	15
Baker et al. (2007)	3
Biondo et al. (2005)	1
Grifoll-Romero et al. (2018)	14
Levitz et al. (2001)	1
Web-blast	55
Similarity search by stand-alone blast ^a	10
PDB information ^b	11
Total	110

^aSimilarity search means that we performed stand-alone blast in the host genome or proteome as a query for each protein (Altschul *et al.*, 1997; Camacho *et al.*, 2009; Sayers *et al.*, 2020).

^bIn the RCSB PDB site (www.rcsb.org), using the plus button in protein feature view, we confirmed the existence of additional protein sequences that shared the same PDB in other organisms.