

A

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270 E P L F T A D F V L A T L D V T N V T Q P T P P P I R D A L
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1011 ATCTTCAAGCGTTGTTTTGAGACCTCTTTTCGCGAGCGGATTCAAAGTAATGAACAGATACTTGCACTTGATTGAGATGAAGATTAATTT
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1101 GTTTTTCTTTACGTTTGTAAACCATGTATGTTTATTAATATCAGCTGTGATCTGAT TTGTCACTAAAAA
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1191 AAAAAAAA

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B

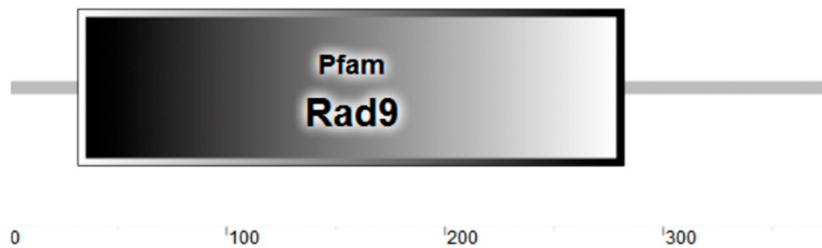


Figure S1. (A) Nucleotide sequence and deduced amino acid sequence of Rad9 gene in *A. sinica*. (B) Results of domain analysis of putative *As-RAD9* protein.

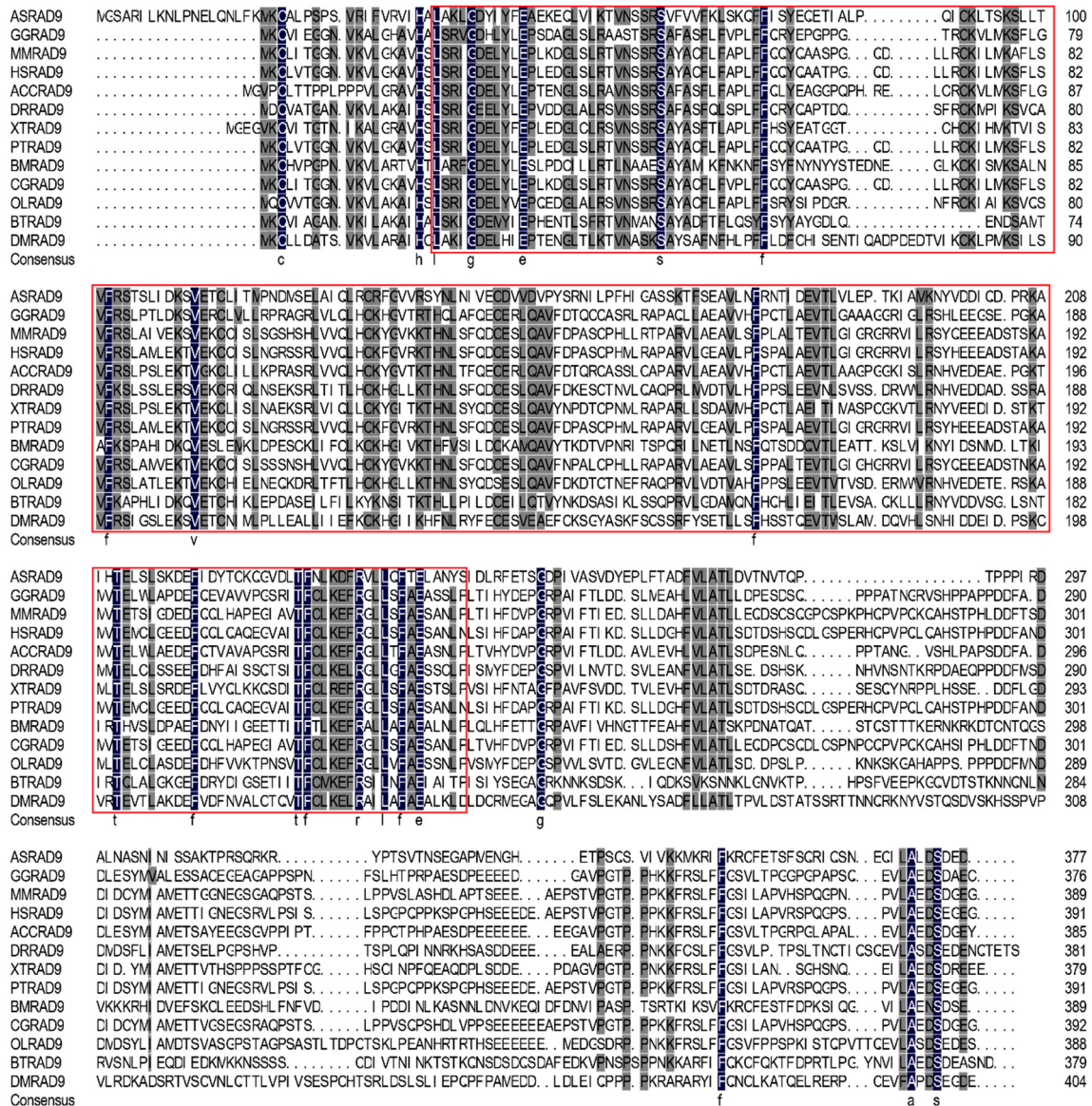


Figure S2: Multiple sequence alignment of *As*-RAD9 protein. Sequence alignment of known RAD9 sequences from 13 species. The sequences and their accession numbers are as follows: MmRAD9, *Mus musculus*, NP_035367.1; CgRAD9, *Cricetulus griseus*, XP_003509986.1; HsRAD9, *Homo sapiens*, NP_004575.1; PtRAD9, *Pan troglodytes*, XP_016776852.1; GgRAD9, *Gallus gallus*, NP_998748.1; AccRAD9, *Aquila chrysaetos canadensis*, XP_011599156.1; XtRAD9, *Xenopus tropicalis*, NP_001005810.1; DrRAD9, *Danio rerio*, NP_956501.2; OIRAD9, *Oryzias latipes*, XP_004073282.1; BmRAD9, *Bombyx mori*, XP_004926904.1; BtRAD9, *Bombus terrestris*, XP_020723628.1; AsRAD9, *Artemia sinica*, (MH_797557); DmRAD9, *Daphnia magna*, KZS14537.1. The sequence of the RAD9 domain is shown in red.

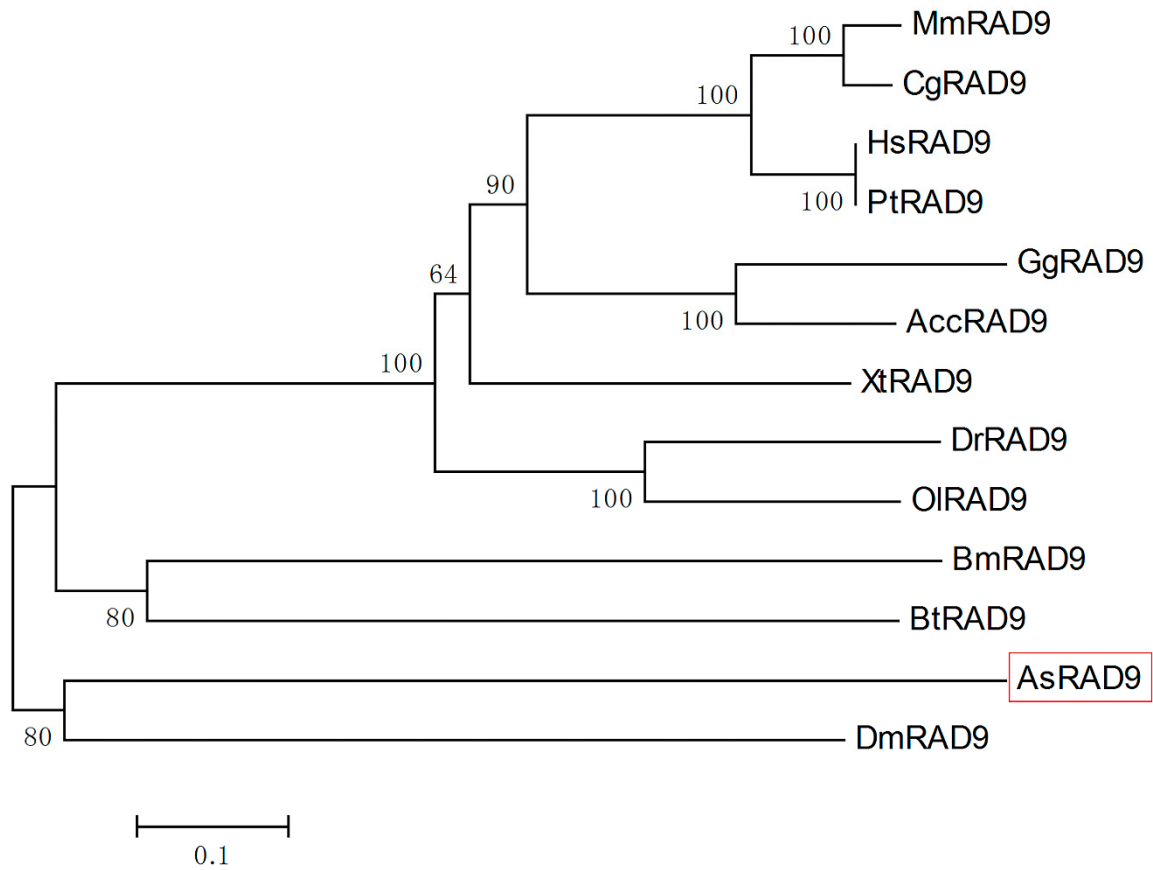


Figure S3: Phylogenetic tree constructed by RAD9 proteins. The sequence and registration number of the RAD9 are the same as in the legend of Figure S2. The red diamond indicates *As*-RAD9 from *A. sinica*.