

**S15 Table. Functional annotation of the genes in the Yap hadal snailfish genome.**

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<b>Parameter</b>	<b>Number</b>	<b>Percentage (%)</b>
Total	24,329	-
Swissprot	22,208	91.28
Nr	23,626	97.11
KEGG	20,200	83.03
InterPro	24,196	99.45
GO	22,460	92.32
PFAM	19,164	78.77
Annotated	24,265	99.74
Unannotated	64	0.26

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