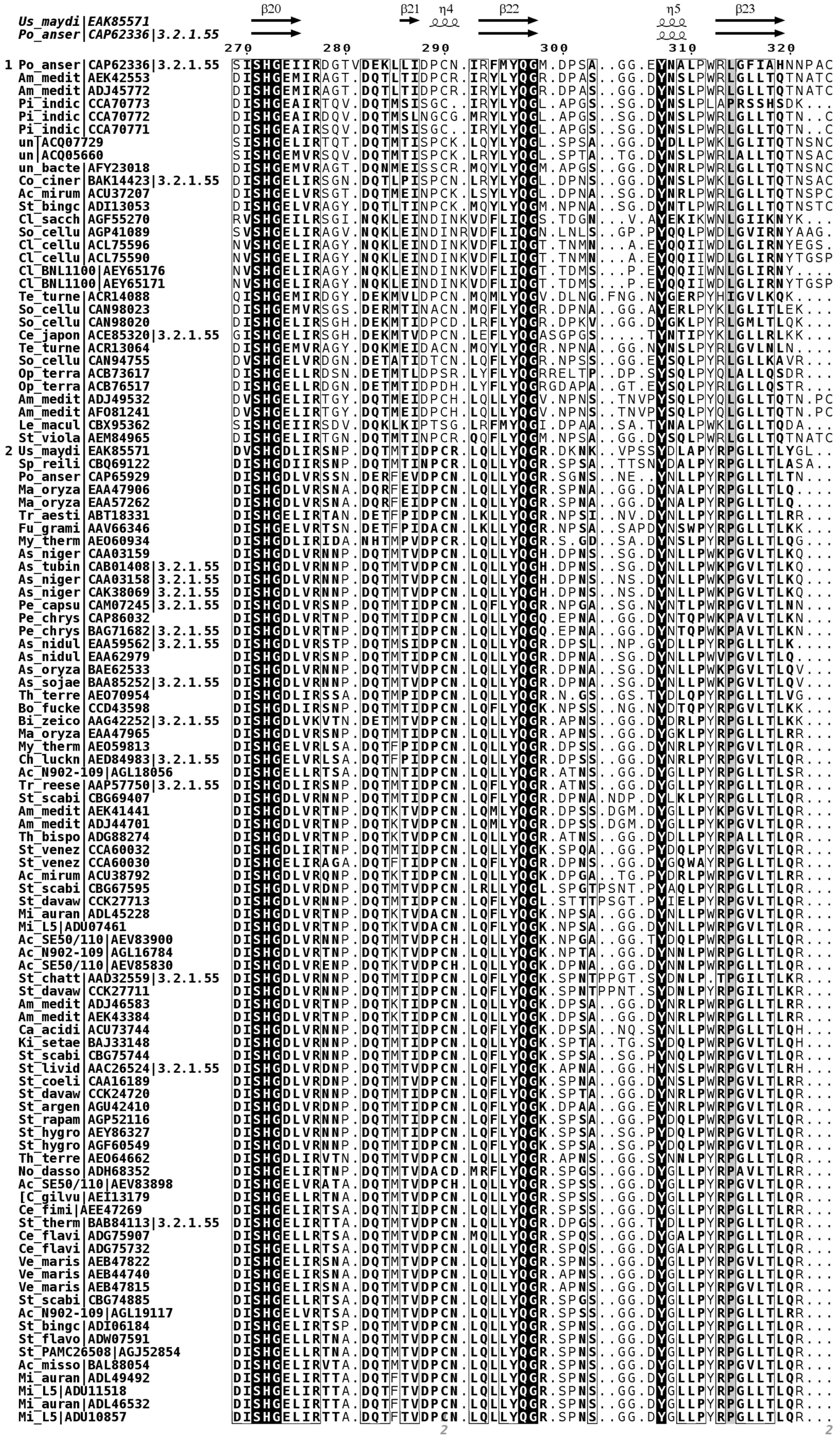
Figure S1

FIGURE S1: Sequence alignment of the GH62 subfamilies. Subfamilies are numbered on the right hand side with the first representative of the family been *Pa*Abf62A and *Um*Abf62A for subfamily 1 and 2, respectively. Secondary structure elements are represented in the top of the page. Um_maydi stand for *Ustilago maydis*, and Po_anser for *Podospora anserina*. Accession numbers are indicated for each protein. pSpript was used to generate this picture (http://espript.ibcp.fr/ESPript/ESPript)(45).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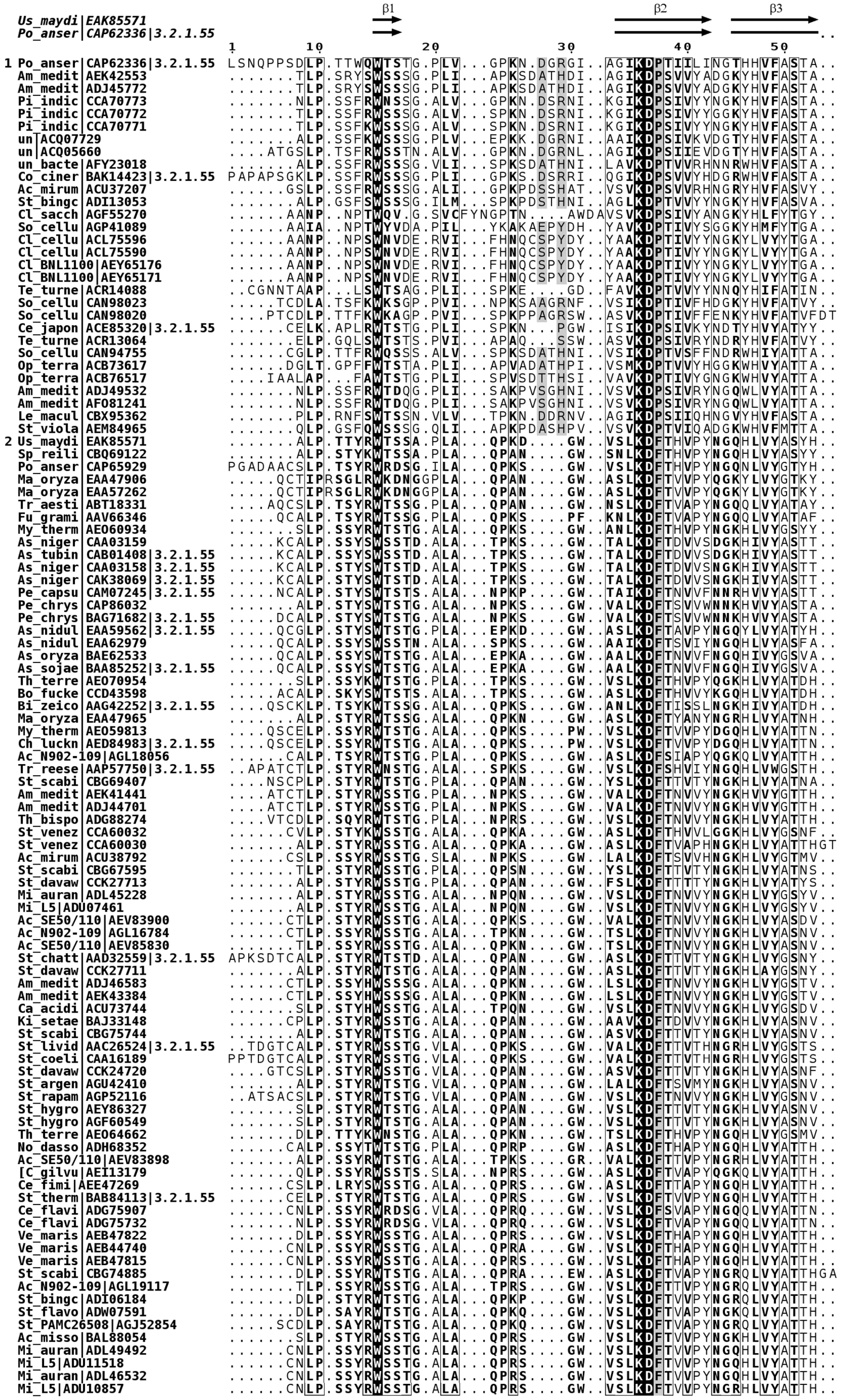
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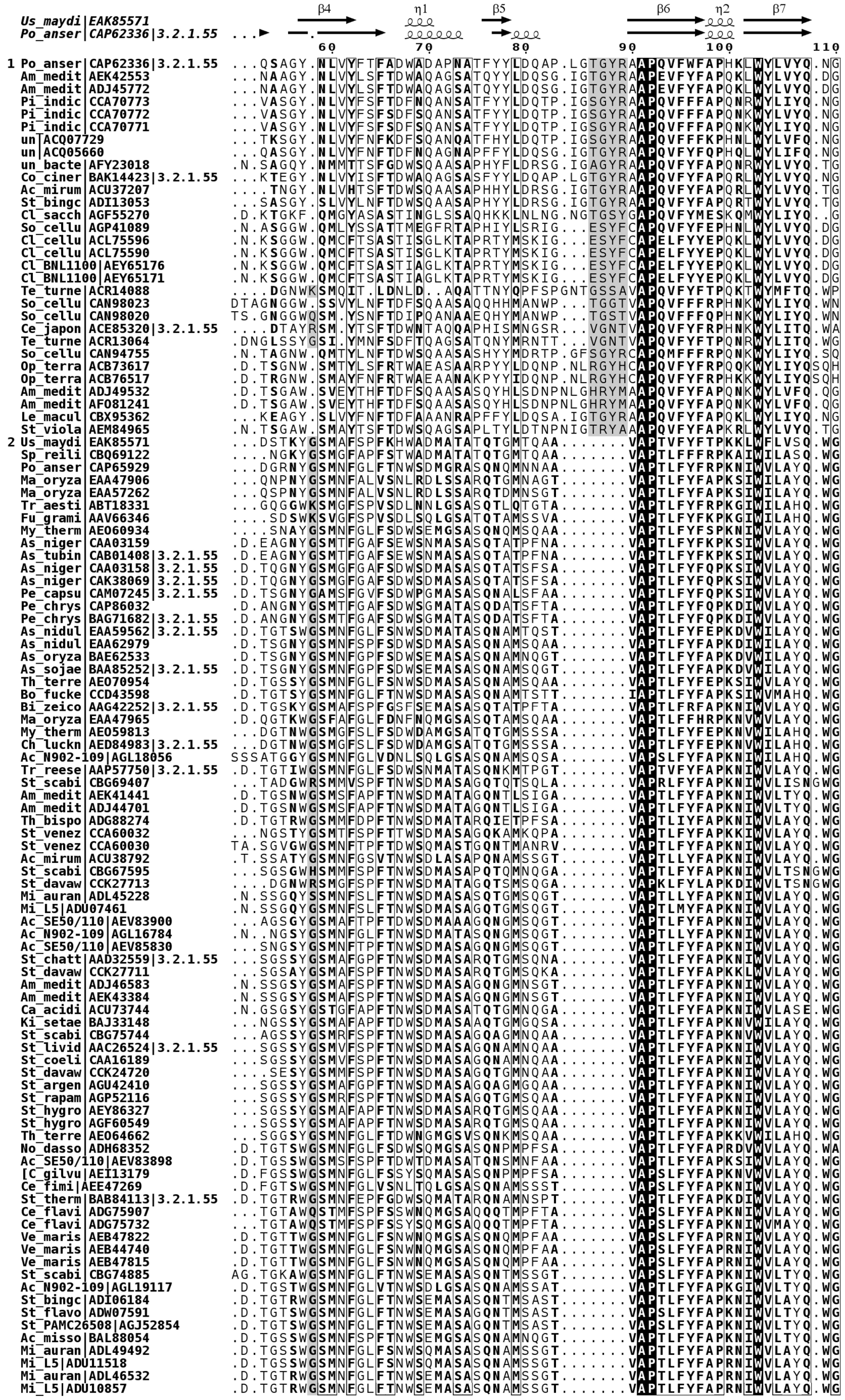
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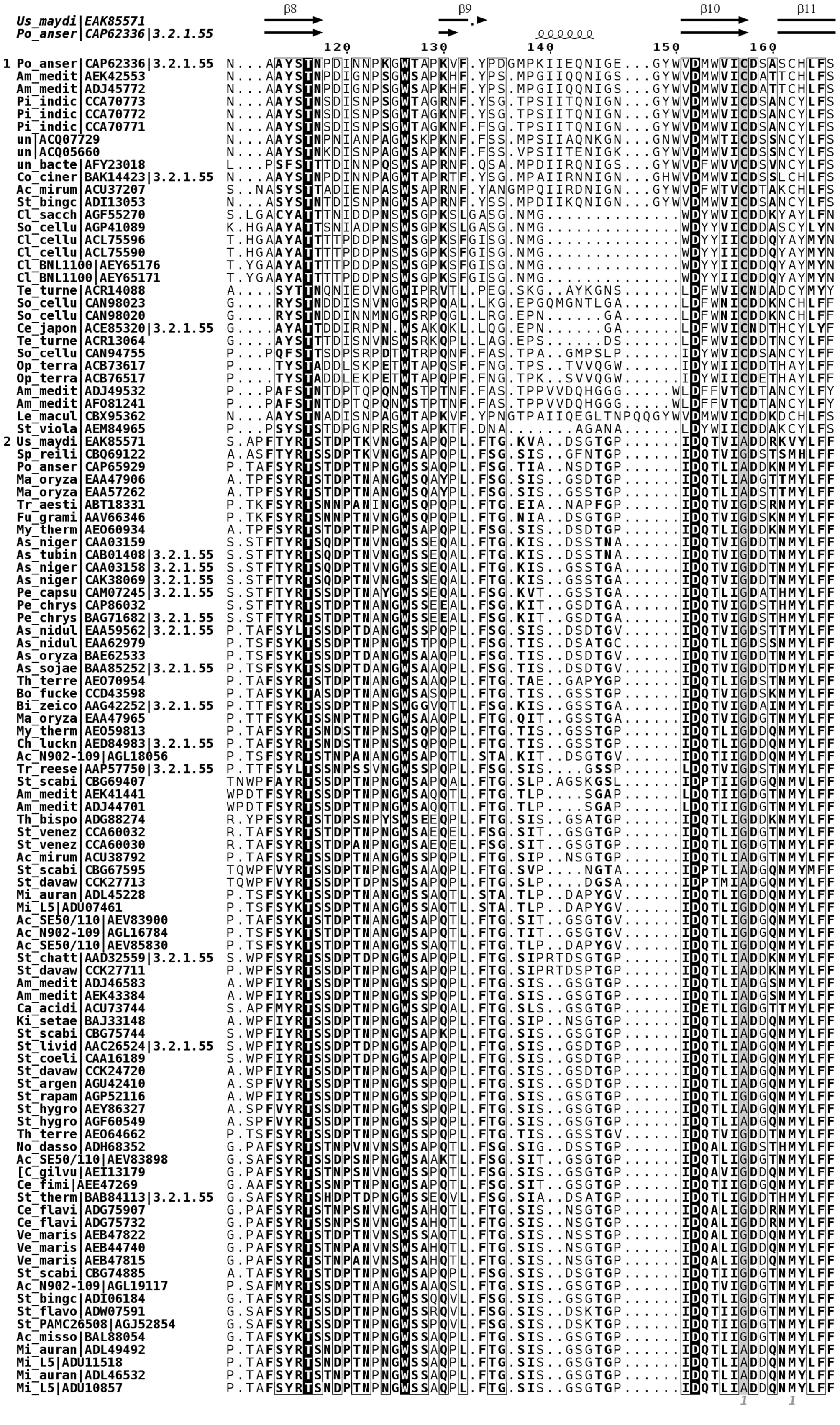
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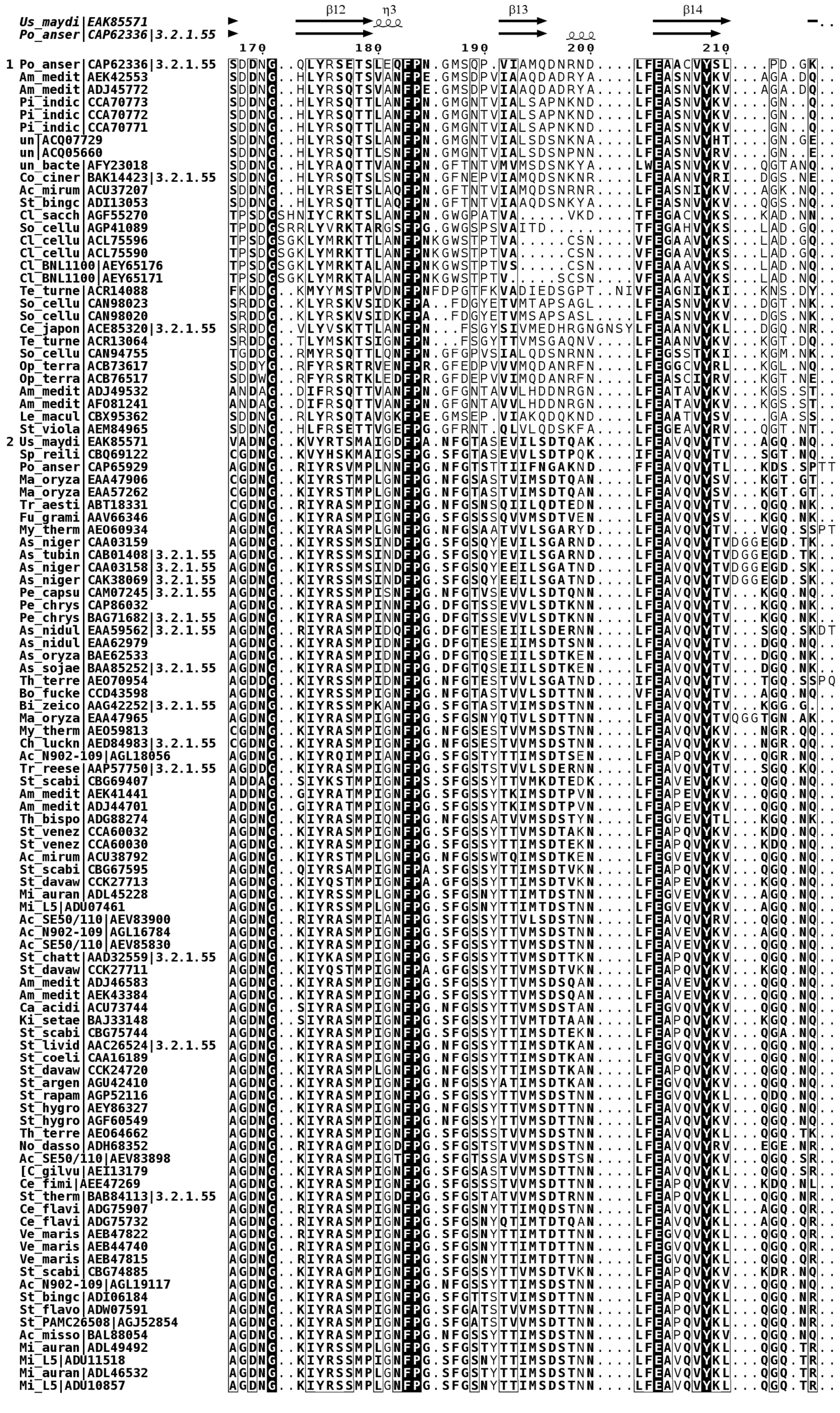

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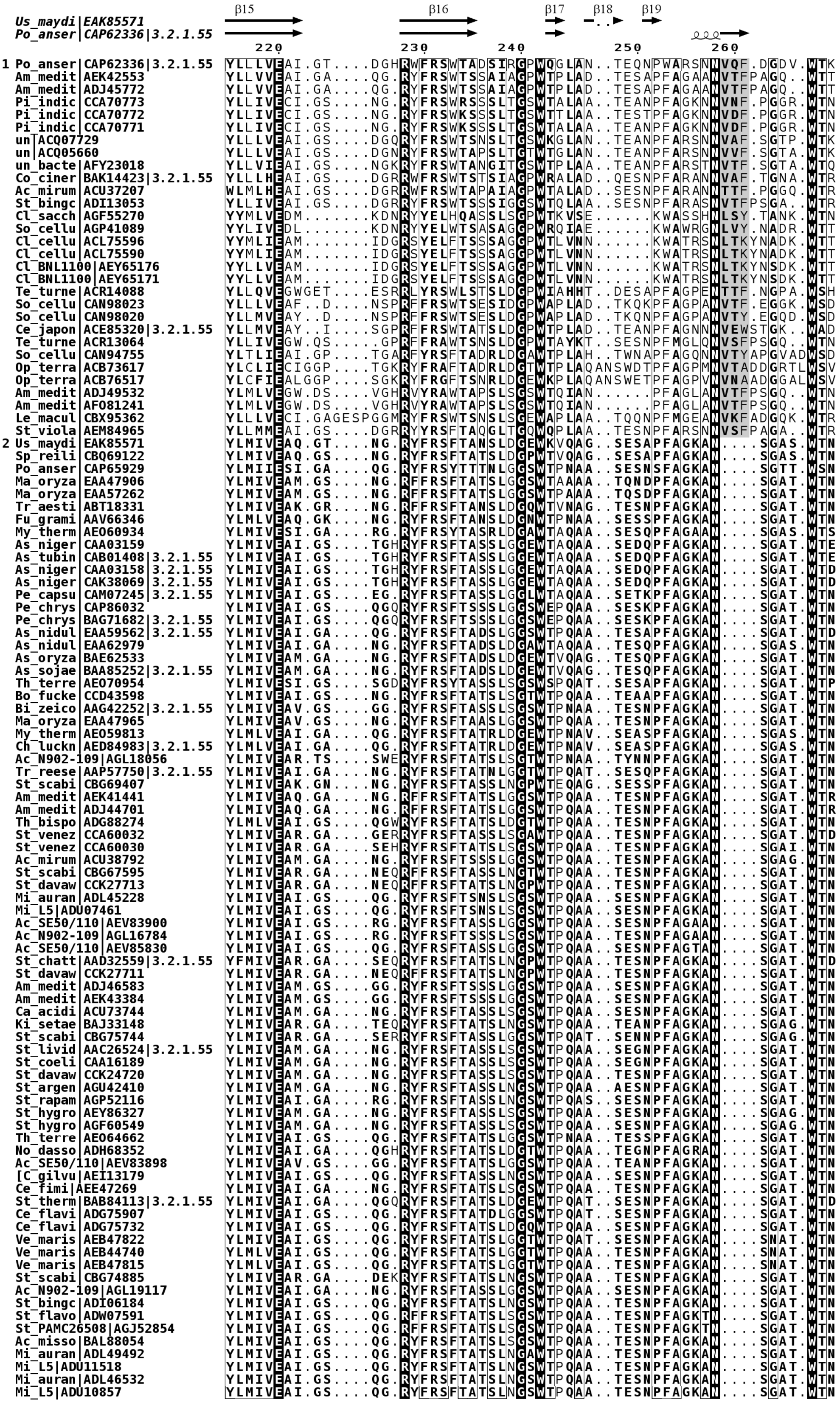
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