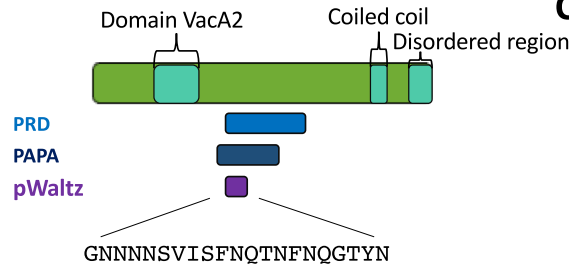


Expanded View Figures

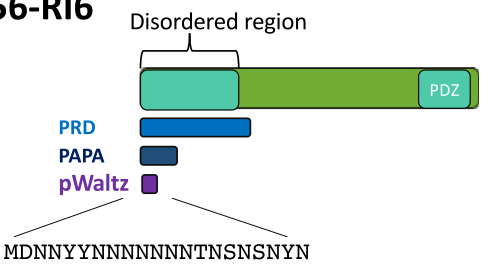
Figure EV1. Sequential information regarding the ten prion-like sequences selected from bacteria found in the gut.

Diagram showing the prion-like regions predicted for all the amyloid cores selected. In green are the main sequences. In light blue are the prion domains (PrD) predicted by PLAAC. In dark blue are the prion domains predicted by PAPA. In purple, the amyloid cores predicted by pWALTZ. Turquoise and yellow are other domains.

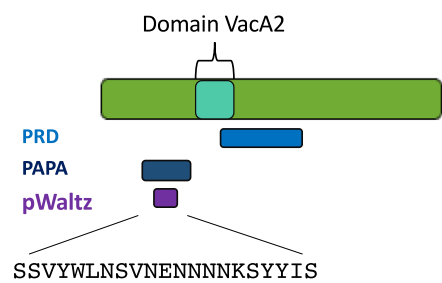
M5YJZ4-HP1



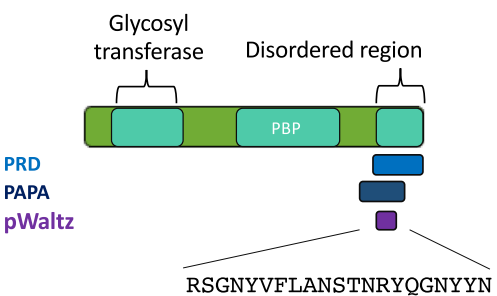
C0FU56-RI6



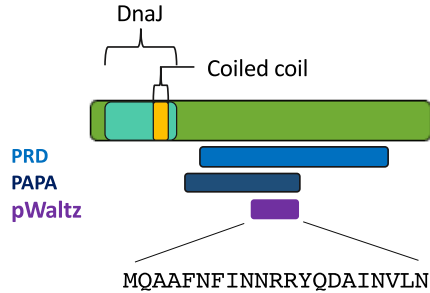
M3SJ19-HP2



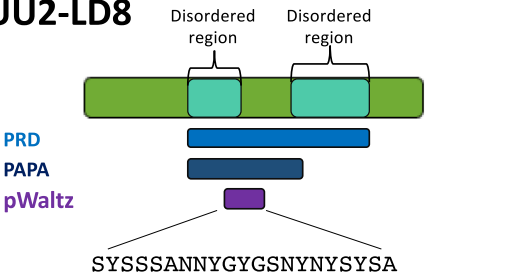
E7GXS5-SA7



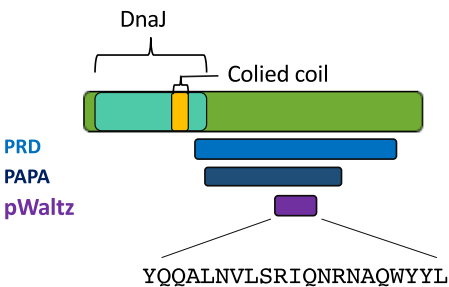
C0B555-CC3



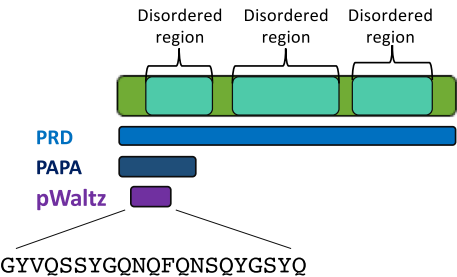
F0HUU2-LD8



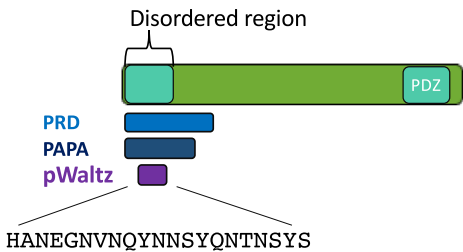
C9L6N5-BH4



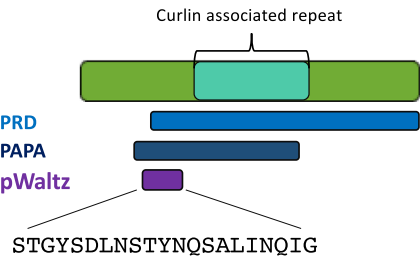
F5LFB6-P9

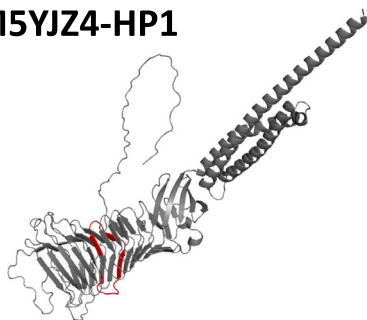
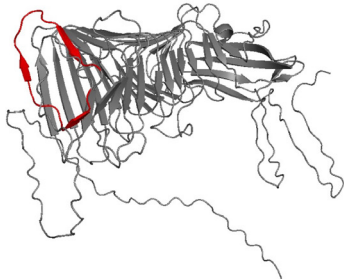
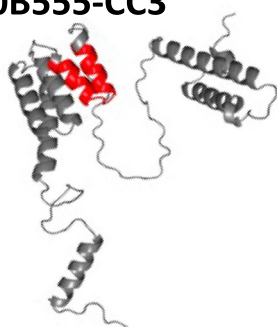
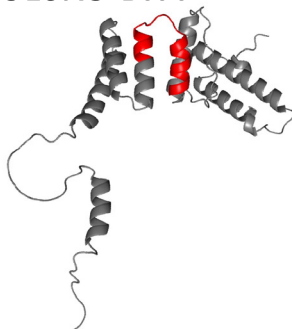
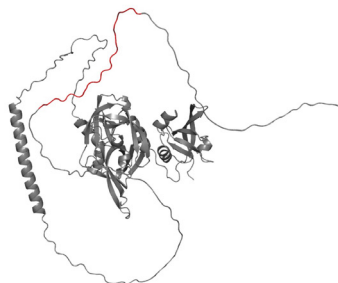
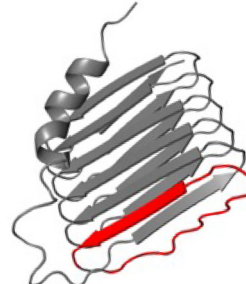


D4KZ46-RI5



G9Y7N7-HA10



M5YJZ4-HP1**C0FU56-RI6****M3SJI9-HP2****E7GXS5-SA7****C0B555-CC3****F0HUU2-LD8****C9L6N5-BH4****F5LFB6-P9****D4KZ46-5RI****G9Y7N7-HA10**

◀ **Figure EV2. Predicted structure of the ten prion-like sequences selected from bacteria found in the gut.**
The images show the Alpha-fold prediction of the whole protein structure.