

Table of protein sequence alignments. The left column contains the identifier 'QUERY' and UniRef90 accession numbers (e.g., UniRef90_F7B4L5, UniRef90_UPI0003C146F2). The right column shows residue positions 1, 10, 20, 30, and 40. The main body of the table displays the amino acid sequences for each query, with residues color-coded: red for conserved residues, yellow for specific substitutions, and black for other residues. Some positions are blank, indicating gaps or missing data. The sequences are aligned across the top, with the UniRef90 identifier at the top left of each row.

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UniRef90_UPI0003EA919F	GRRRIKPAVILKYL	LTNNHSGFRFSEELKPPGGCLKVVML	LL
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UniRef90_W5UKC0	GRRRIKPAVILKYL	LTNNHSGFRFSEELKPPGGCLKVVML	LL
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