

logo

NTDB id 165105 MGAS27061 RS01320 WP 024623426.1  
NTDB id 475 HSISS4 RS07005 WP 021143809.1  
NTDB id 324 STU RS16140 WP 011226306.1  
NTDB id 292 STER RS06940 WP 011681419.1  
consensus



80  
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76

logo

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consensus



159  
156  
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156

logo

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consensus



238  
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236

logo

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consensus



318  
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316

logo

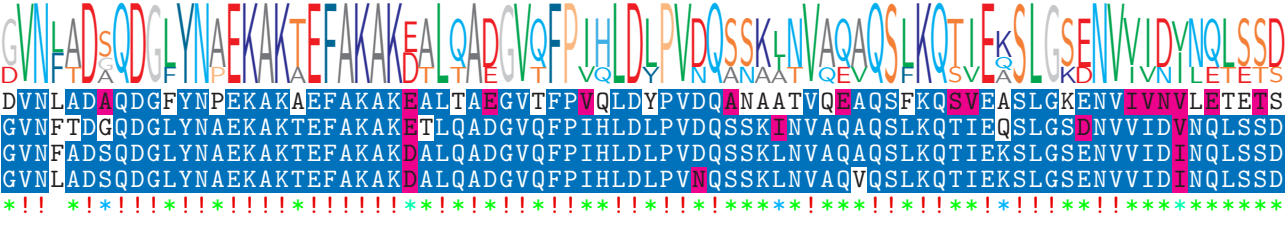
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397  
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396

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477  
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476

logo

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consensus



Sequence logo showing amino acid conservation across four species. The y-axis represents information content. The x-axis shows positions 1 to 20. The consensus sequence is: **THQAQGFYIETPEQQYDIISSWCGPDYQDPRTYLDIMSPVGGGSVIQKLGIKAGQNKDVVAAAAGLDTYQTLLDEAAAI**

556

555

555

555

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consensus



Sequence logo showing amino acid conservation across four species. The y-axis represents information content. The x-axis shows positions 1 to 20. The consensus sequence is: **TSDLNVRVDRYAQAQAWLEDSSLVPLTVGNGAAPVISRLTPFTGASLQVGDKNSSNDYIKYVKPQEKVTKKEYEQSRE**

633

634

635

635

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consensus



Sequence logo showing amino acid conservation across four species. The y-axis represents information content. The x-axis shows positions 1 to 20. The consensus sequence is: **KWLKEKKESNEKAQKDLEKHVK**

656

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657

-  non conserved
-  similar
-  ≥ 50% conserved