

			10	20	30	40	
<i>ARSA_MOUSE</i>	409	S D P A C H A - A N R L T A H E P P L L Y D L S Q D P G E N Y N V L E S I E G V S P E A					451
<i>Q9DC66_MOUSE</i>	409	S D P A C H A - A N R L T A H E P P L L Y D L S Q D P G E N Y N V L E S I E G V S P E A					451
<i>Q32K15_CANFA</i>	410	P D P A C H A - S S P L T A H E P P L L F D L S E D P G E N Y N L L G G M A E V A P E A					452
<i>Q5BL32_BRARE</i>	409	P D N S C S L - L A F L K Y H D P P L L F N L E T D P S E N Y N L D G D - - - - - Q W					445
<i>Q6AX40_XENLA</i>	410	P D P D C H V - T A L L K S H D P P L L F D L S T D P A E N Y N L L K D G - - I P V D L					450
<i>Q8WNR3_PIG</i>	410	A D P A C H A - S S P L T A H E P P L L F D L S E D P G E N Y N L L G G V A E V A P E V					452
<i>Q5XFU5_MOUSE</i>	436	G A K A C D G S V G P E Q H H V A P L I F N L E D A A D E G M P L Q K - - - - G S P E Y					475
<i>Q3TYD4_MOUSE</i>	436	G A K A C D G S V G P E Q H H V A P L I F N L E D A A D E G M P L Q K - - - - G S P E Y					475
<i>ARSG_HUMAN</i>	436	G A R A C D G S T G P E L Q H K F P L I F N L E D D T A E A V P L E R - - - - G G A E Y					475
<i>Q32KJ9_RAT</i>	436	G A K A C D G G V G P E Q H H V S P L I F N L E D D A A E S S P L Q K - - - - G S P E Y					475
Consensus		G D P A C H + S V G P L T A H E P P L L F + L + D D P G E N Y N L L K + I + + + S P E Y					