

Figure 1 figure Supplement 3

ICE1 Predicted MIF4G Domain

Figure 1 displays the multiple sequence alignment of the C-terminal region of the protein across six species: Human, Rat, Mouse, Bat, Chicken, Killifish, Zebrafish, and Fruitfly. The alignment is shown in three blocks, each with a scale bar at the bottom. The scale bars indicate the percentage of identity between the sequences: 100% (black), 90% (dark grey), 80% (medium grey), 70% (light grey), 60% (very light grey), 50% (white), 40% (lightest grey), 30% (lightest grey), 20% (lightest grey), 10% (lightest grey), 0% (white). The alignment shows high conservation across all species, with the C-terminal region being particularly conserved.