

cubitus interruptus

Sequence alignment for cubitus interruptus across species: Er\_ci, Ek\_ci, Tc\_ci, Al\_ci, Gm\_ci, Dm\_ci. Includes amino acid sequences and residue numbers (20, 40, 60, 80, 100, 120, 140, 160, 180, 200).

engrailed

Sequence alignment for engrailed across species: Er\_en, Ek\_en, Gm\_en, Af\_en, Tc\_en, Pd\_en, Al\_en. Includes amino acid sequences and residue numbers (20, 40, 60, 80, 100, 120, 140, 160, 180, 200, 220, 240, 260, 280, 300, 320, 340, 360, 380, 400, 420, 440).

hedgehog

Sequence alignment for hedgehog across species: Er\_hh, Ek\_hh, Gm\_hh, Af\_hh, Tc\_hh, Pd\_hh. Includes amino acid sequences and residue numbers (20, 40, 60, 80, 100, 120, 140, 160, 180, 200, 220, 240, 260, 280, 300, 320, 340, 360, 380, 400, 420, 440, 460, 480, 500, 520, 540).

wingless

Sequence alignment for wingless across species: Er\_wg, Ek\_wg, Ap\_wg, Dm\_wg, Dp\_Wnt1, Tc\_wg, Al\_wg, Is\_wg, Gm\_Wnt1, Ct\_Wnt1, Lg\_Wnt1, Pd\_Wnt1, Hs\_Wnt1, Hr\_Wnt1, Nv\_Wnt1, Dm\_Wnt6, Gm\_Wnt6, Tc\_Wnt6, Nv\_Wnt6. Includes amino acid sequences and residue numbers (20, 40, 60, 80, 100, 120, 140, 160, 180, 200, 220, 240, 260, 280, 300, 320, 340, 360, 380, 400, 420).