



Figures 6-1: Generalised secondary structure of the tRNA<sup>Leu</sup> (UAA) in black and the *trn*<sub>UAA</sub> group I intron in orange (redrawn from KUHSEL *et al.* 1990, XU *et al.* 1990). Note the UAA-anticodon (yellow) of the tRNA (green), pairing regions 1 through 9 (8 being very long) and conserved motifs P, Q, R, and S (red).