Instructions for Building the Paper Model of tRNA



Additional information about this activity can be found at: bit.ly/HVzIoH www.rcsb.org • info@rcsb.org



Acceptor arm D-arm D-arm Anticodon Anticodon



5 Fold in ("valley fold") on the small vertical dotted lines.



6 Bring colored/patterned sections together, and tape so the grey tab is hidden. It is important that you match the bases in the following order:



 1. 10-13 with 25-22
 0

 2. 27-31 with 43-39
 0

 3. 49-53 with 65-61
 0

 4. 1-7 with 72-66
 0



Notice the almost-perfect base pairing (G:C and A:U bases pair up). At this point the model is a cloverleaf shapethe secondary structure of tRNA. Each colored region represents the double helical regions of the structure.

Acceptor Arm

Anticodon Loop



III. The tertiary structure



Bring together bases G19 and C56 (colored yellow) to form a base pair – hide the grey tab by taping it under. This forms the beginnings of the tRNA tertiary structure– the inverted L-shape.

To further explore details of tertiary structure see the atomic model of tRNA at **bit.ly/HVzIoH**

What is tRNA?

Transfer RNA (tRNA)¹ "translates" the genetic code into the language of proteins. Each tRNA molecule binds to a specific amino acid on the acceptor arm, recognizes its corresponding code on the mRNA through the anticodon loop region and delivers the amino acid to a growing peptide chain in the ribosome² for protein synthesis.

1. See the *Molecule of the Month* feature at doi: 10.2210/rcsb_pdb/mom_2001_3

2. See the Molecule of the Month feature at doi: 10.2210/rcsb_pdb/mom_2010_1







