RNAfold and forna input parameters for sequence structures

RNAfold:

* Fold algorithms and basic options
  + Minimum free energy and partition function – checked
  + Avoid isolated base pairs – unchecked
* Energy parameters
  + DNA parameters (Mathews model, 2004)
  + Temperature – 22 C
  + Salt concentration – 1.021 M

Forna:

* Colour scheme
  + Custom
    - Full sequences:
      * Input: 13-14, 19-21, 26-28, 46-48, 54-56, 61-62:lightgreen
    - IL6-4202.1 truncate:
      * Input: 4-5, 10-12, 17-19:lightgreen
    - IL6-7326.1 truncate: 2-3, 8-10, 15-17:lightgreen