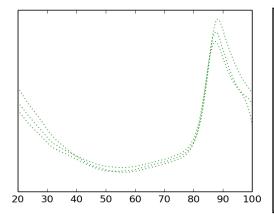
MELTDOWN v2.0.0 Melt Curve Analysis

apo-DHDPS-perugini_buffer screen 9_010714...





Full interpretation of the results requires you to look at the individual melt curves.

The summary graph appears to be unreliable

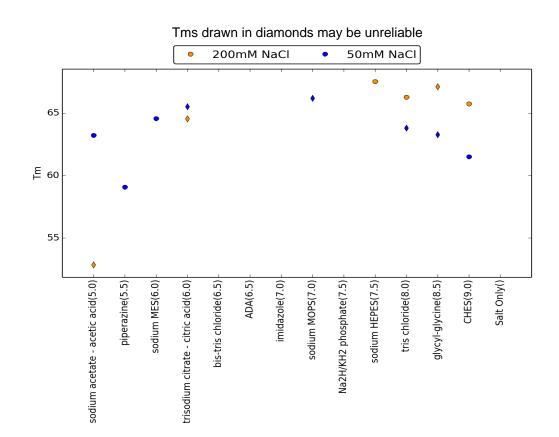
32% of curves were used in Tm estimations

Average estimation of error is 0.2 C

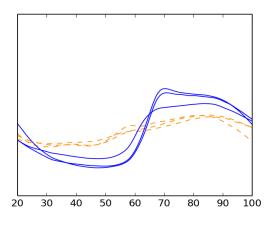
Protein as supplied is not well behaved

Protein as supplied: Tm = N/A

Lysozyme Control: Passed No Dye Control: Passed No Protein Control: Passed



Highest Tm = 67.54 +/- 0.01 (sodium HEPES / 200mM NaCl)



sodium acetate - acetic acid (5.0)

Grouped by Tm

200mM NaCl 52.85 (+/-0.27)^ 50mM NaCl 63.22 (+/-2.06)

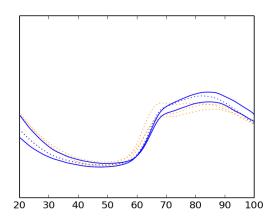
20 30 50 60 70 90 100 80

piperazine (5.5)

Grouped by Adjusted pH at Tm Tm

200mM NaCl None

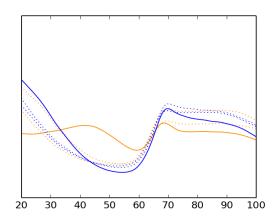
50mM NaCl 59.07 (+/-0.02) 4.79



sodium MES (6.0)

Grouped by Tm Adjusted pH at Tm 200mM NaCl None

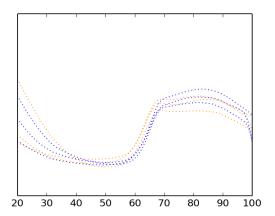
50mM NaCl 64.58 (+/-0.11) 5.65



trisodium citrate - citric acid (6.0)

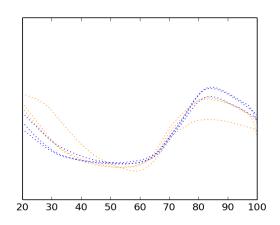
Grouped by Tm Adjusted pH at Tm

200mM NaCl 64.55 6.17 50mM NaCl 65.53 6.18



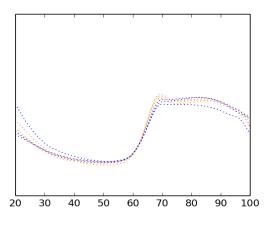
bis-tris chloride (6.5)

Grouped by Tm 200mM NaCl None 50mM NaCl None



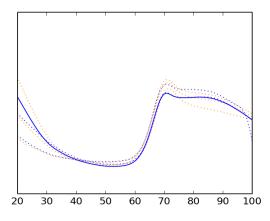
ADA (6.5)

Grouped by Tm 200mM NaCl None 50mM NaCl None



imidazole (7.0)

Grouped by Tm
200mM NaCl None
50mM NaCl None

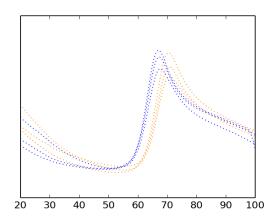


sodium MOPS (7.0)

Grouped by Tm Adjusted pH at Tm

200mM NaCl None

50mM NaCl 66.19 6.62

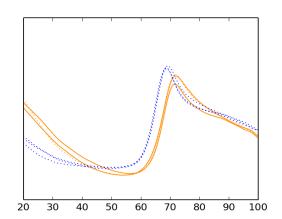


Na2H/KH2 phosphate (7.5)

Grouped by Tm

200mM NaCl None

50mM NaCl None

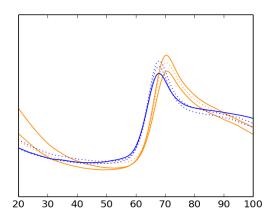


sodium HEPES (7.5)

Grouped by Tm Adjusted pH at Tm

200mM NaCl 67.54 (+/-0.01) 7.21

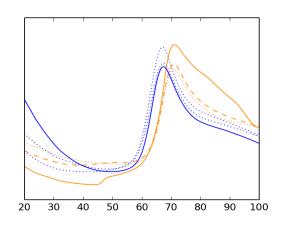
50mM NaCl None



tris chloride (8.0)

Grouped by Tm Adjusted pH at Tm 200mM NaCl 66.29 (+/-0.15) 7.06

50mM NaCl 63.82 7.11



glycyl-glycine (8.5)

Grouped by Tm Adjusted pH at Tm 200mM NaCl 67.12 (+/-0.28)^ 7.71

50mM NaCl 63.27 7.78

