

```

(*)
holo = %holo im Gleichgewicht
b= Kpre= [apoA]/[ApoB]
c= logKd
l= Anfangskonzentration von Adenine
r= Anfangskonzentration von RNA
*)
R = 8.3144621 (*J/mol*K*)
DeltaHbind = 110470.2357608338`
DeltaSbind = 238.303284932985`
DeltaHpre = 52002.80824997664`
DeltaSpre = 167.28088132812806`
(*Daten sind aus Kd_und_Kpre_function_of_T.nb the
values for Kd and Kpre were taken from the Nature xls list.*)
holo[T_, r_, l_] :=

```

$$\left(100 * \left(\left(\frac{1}{2 e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}}} \right) \left(10^{\text{Log}\left[10, e^{-\frac{(\text{DeltaHbind}-T*\text{DeltaSbind})}{R*T}}\right]} + 10^{\text{Log}\left[10, e^{-\frac{(\text{DeltaHbind}-T*\text{DeltaSbind})}{R*T}}\right]} \right) e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}} + \right. \\
 \left. e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}} l + e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}} r - \right. \\
 \left. \sqrt{\left(-4 \left(e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}} \right)^2 l r + \left(-10^{\text{Log}\left[10, e^{-\frac{(\text{DeltaHbind}-T*\text{DeltaSbind})}{R*T}}\right]} - 10^{\text{Log}\left[10, e^{-\frac{(\text{DeltaHbind}-T*\text{DeltaSbind})}{R*T}}\right]} \right) \right.} \\
 \left. \left. e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}} - e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}} l - e^{-\frac{(\text{DeltaHpre}-T*\text{DeltaSpre})}{R*T}} r \right)^2 \right) \right) / r \Big)$$

8.31446

110470.

238.303

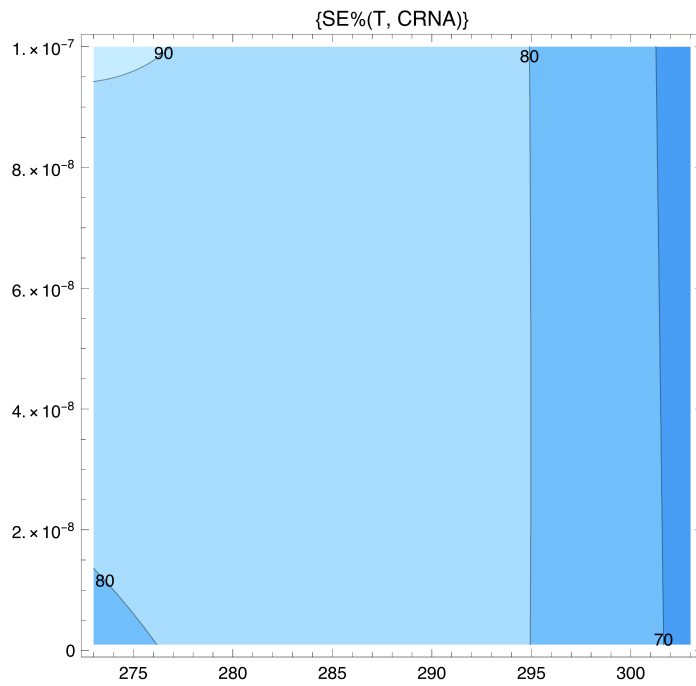
52002.8

167.281

```

p1 = ContourPlot[holo[T, r, 1.5 * 10-6] - holo[T, r, 0.01 * 10-6], {T, 273, 303},
  {r, 1 * 10-9, 100 * 10-9}, Contours → {10, 20, 30, 40, 50, 60, 70, 80, 90, 100},
  ColorFunction → ColorData["DeepSeaColors"], ContourLabels → True,
  PlotLabel → {"SE%(T, CRNA)"}, PlotRange → {0, 100}]

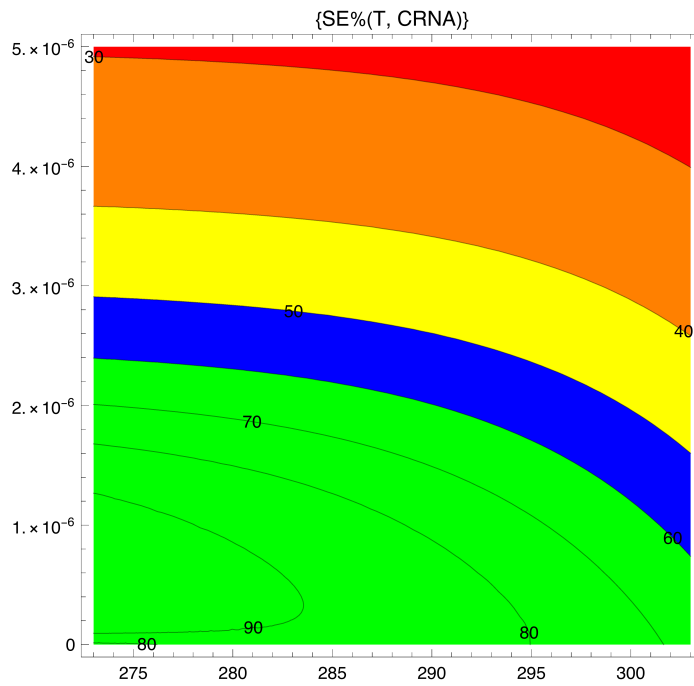
```



```

p1 = ContourPlot[holo[T, r, 1.5 * 10-6] - holo[T, r, 0.01 * 10-6], {T, 273, 303},
  {r, 1 * 10-9, 5 * 10-6}, Contours → {10, 20, 30, 40, 50, 60, 70, 80, 90, 100},
  ContourShading → {Red, Red, Red, Orange, Yellow, Blue, Green, Green, Green, Green},
  ContourLabels → True, PlotLabel → {"SE%(T, CRNA)"}, PlotRange → {0, 100}]

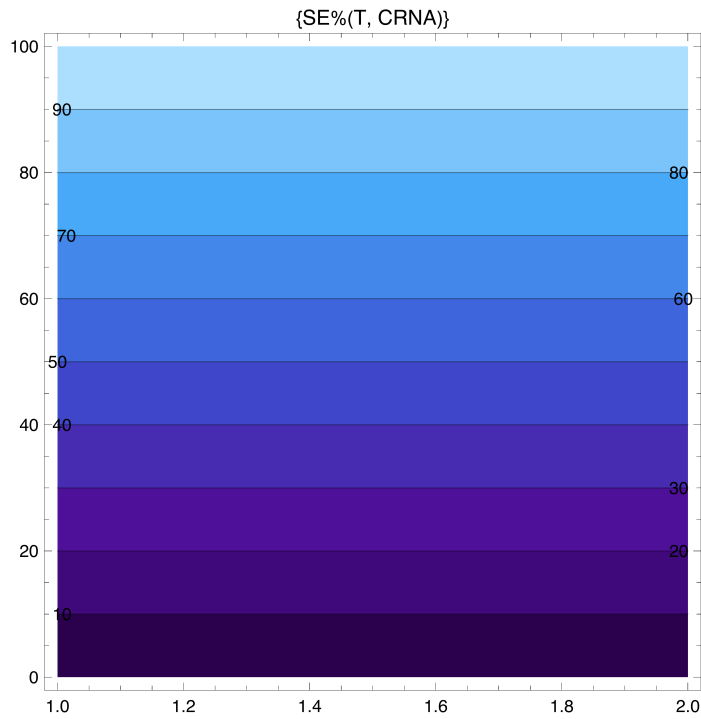
```



```

schoen[x_, y_] := (x / x) + y - 1
p1 = ContourPlot[schoen[x, y], {x, 1, 2},
  {y, 0, 100}, Contours → {10, 20, 30, 40, 50, 60, 70, 80, 90, 100},
  ColorFunction → ColorData["DeepSeaColors"], ContourLabels → True,
  PlotLabel → {"SE%(T, CRNA)"}, PlotRange → {0, 100}]

```



```

150 * 106 / (30 * 60 * 24 * 7 * 4 * 12) // N

```

```

10.334

```