

```
[ > restart;
  VLE data for chloroform(1)/1,4-dioxane(2) at 50 degC (S,vN&A Table 11.3)
```

## Input data

```
[ > P :=
  [15.79,17.51,18.15,19.30,19.89,21.37,24.95,29.82,34.80,42.10,60
  .38,65.39,69.36];
  P :=
  [15.79, 17.51, 18.15, 19.30, 19.89, 21.37, 24.95, 29.82, 34.80, 42.10, 60.38, 65.39, 69.36]
[ > x1 :=
  [0.0,.0932,.1248,.1757,.2,.2626,.3615,.4750,.5555,.6718,.878,.9
  398,1.0];
  x1 := [0, .0932, .1248, .1757, .2, .2626, .3615, .4750, .5555, .6718, .878, .9398, 1.0]
[ > y1 :=
  [0.0,.1794,.2383,.3302,.3691,.4628,.6184,.7552,.8378,.9137,.986
  0,.9945,1.0];
  y1 := [0, .1794, .2383, .3302, .3691, .4628, .6184, .7552, .8378, .9137, .9860, .9945, 1.0]
[ > with(linalg): with(plots):with(stats):
  Warning, new definition for norm
  Warning, new definition for trace
[ > n := coldim([x1]);
  n := 13
[ > plotdata := (i0,i1,x,y) -> plot([[x['i'],y['i']]
  $'i'=i0..i1],color=black,style=point,symbol=circle);
  plotdata :=
  (i0, i1, x, y) → plot([[xi, yi] $ ('i' = i0 .. i1)], color = black, style = point, symbol = circle)
[ > Pvsx := plotdata(1,n,x1,P):
  Pvsy := plotdata(1,n,y1,P):
  display({Pvsx,Pvsy});
```

## Very crude minimization routine for later use

Finds rough minimum of a function of two variables. Examines value of function over an equally-spaced grid of values of the two variables. "fn" is the function, which should take two arguments; "nPts" is the number of grid points for each variable; "x1Range" and "x2Range" are lists (of the form [x1min,x1max]) which specify the upper and lower bounds of the grid for each variable. You can refine the search by running the routine several times, each one with a narrower range of values for the two variables.

```
[ > crudeMinimize := proc(fn,nPts,x1Range,x2Range)
  local x1,x2,m1,m2,x1step,x2step,obj,bestobj,x1best,x2best;
  x1step := (x1Range[2]-x1Range[1])/nPts;
  x2step := (x2Range[2]-x2Range[1])/nPts;
  bestobj := 1e32;
  for m1 from 0 to nPts do
    x1 := x1Range[1] + m1*x1step;
    for m2 from 0 to nPts do
      x2 := x2Range[1] + m2*x2step;
      obj := fn(x1,x2);
      if(obj < bestobj) then
        bestobj := obj;
```

```

                x1best := x1;
                x2best := x2;
            fi;
        od;
    od;
    evalf([x1best,x2best,bestobj]);
end;
crudeMinimize := proc(fn, nPts, x1Range, x2Range)
local x1, x2, m1, m2, x1step, x2step, obj, bestobj, x1best, x2best;
    x1step := (x1Range[2] - x1Range[1]) / nPts;
    x2step := (x2Range[2] - x2Range[1]) / nPts;
    bestobj := .1*10^33;
    for m1 from 0 to nPts do
        x1 := x1Range[1] + m1*x1step;
        for m2 from 0 to nPts do
            x2 := x2Range[1] + m2*x2step;
            obj := fn(x1, x2);
            if obj < bestobj then bestobj := obj; x1best := x1; x2best := x2 fi
        od
    od;
    evalf([x1best, x2best, bestobj])
end

```

## Examine and fit Gibbs excess

### Determine gE from data

```

> p2sat := 15.79;  p1sat := 69.36;
                                p2sat := 15.79
                                p1sat := 69.36
> x2 := [seq(1-x1[i],i=1..n)]:
y2 := [seq(1-y1[i],i=1..n)]:
gamma1 := [gamlinf,seq(P[i]*y1[i]/x1[i]/p1sat,i=2..n-1),1]:
gamma2 := [1,seq(P[i]*y2[i]/x2[i]/p2sat,i=2..n-1),gam2inf]:
lngamma1 := [seq(ln(gamma1[i]),i=1..n)]:
lngamma2 := [seq(ln(gamma2[i]),i=1..n)]:
gE := [seq(x1[i]*lngamma1[i] + x2[i]*lngamma2[i],i=1..n)]:
> gEdataPlot := plotdata(1,n,x1,gE):
> display(gEdataPlot);

```

### Fit data to 1-constant and 2-constant (Eq. 11.7) Margules forms

```

1-constant form
gEfit1 := fit[leastsquare[[X,Y], Y=a*X*(1-X)]([x1,gE]);
                                gEfit1 := Y = -0.9615256100 X (1 - X)
> a := -0.9615;
                                a := -0.9615
2-constant form
> gEfit2 := fit[leastsquare[[X,Y], Y=(A21*X +
                                A12*(1-X))*X*(1-X)]([x1,gE]);

```

$$gEfit2 := Y = (-.5084474687 X - .7402510613) X (1 - X)$$

```

gamma2vanLaar := x -> exp(A21p/(1 + A21p*(1-x)/A12p/x)^2);
                                     
$$\left( \frac{A21p}{1 + \frac{A21p(1-x)}{A12p x}} \right)^2$$

                                     gamma2vanLaar := x → e
> gammavanLaarPlots :=
  plot({gammalvanLaar(x), gamma2vanLaar(x)}, x=0..1):
> display({gammaldataPlot, gamma2dataPlot, gammavanLaarPlots});

```

## Fit to Wilson equation

Formula is not linear in the fitting parameters L12 and L21. Must work with numerical minimization routine (define above).

```

> gEWilson := (x, L12, L21) -> -x*ln(x + (1-x)*L12) -
  (1-x)*ln((1-x) + x*L21);
  gEWilson := (x, L12, L21) → -x ln(x + (1-x) L12) - (1-x) ln(1-x + x L21)
> gam1inf := gammalMargules2(0, a12, a21); gam2inf :=
  gamma2Margules2(1, a12, a21);
  gam1inf := .4769708028
  gam2inf := .2868774953

```

Determine initial guess values for Lambda12 and Lambda21 by matching infinite-dilution activity coefficients

```

> solve({ln(gam1inf)=-ln('L12')+1-'L21', ln(gam2inf)=-ln('L21')+
  1-'L12'}, {'L12', 'L21'});
  {L12 = .0004390265140, L21 = 9.471250750}
> l12 := 0.000439; l21 := 9.4713;
  l12 := .000439
  l21 := 9.4713

```

Define functions that give activity coefficients as a function of mole fraction and parameter values

```

> gammalWilson := (x, L12, L21) -> exp(-ln(x + (1-x)*L12) +
  (1-x)*(L12/(x + (1-x)*L12) - L21/((1-x) + x*L21)));
  gamma2Wilson := (x, L12, L21) -> exp(-ln((1-x) + x*L21) -
  x*(L12/(x + (1-x)*L12) - L21/((1-x) + x*L21)));
  gammalWilson := (x, L12, L21) → e 
$$\left( -\ln(x + (1-x)L12) + (1-x) \left( \frac{L12}{x + (1-x)L12} - \frac{L21}{1-x+xL21} \right) \right)$$

  gamma2Wilson := (x, L12, L21) → e 
$$\left( -\ln(1-x+xL21) - x \left( \frac{L12}{x + (1-x)L12} - \frac{L21}{1-x+xL21} \right) \right)$$


```

Check that functions match infinite-dilution values

```

> gammalWilson(0, l12, l21);
  .4769761187
> gamma2Wilson(1, l12, l21);
  .2868760113

```

Check to see how well these coefficients reproduce the gE and activity-coefficient data

```

> gEWilsonPlot := plot(gEWilson(x, l12, l21), x=0..1):
> display({gEWilsonPlot, gEdataPlot});
> gammaWilsonPlots :=
  plot({gammalWilson(x, l12, l21), gamma2Wilson(x, l12, l21)}, x=0..0
  .99):
> display({gammaldataPlot, gamma2dataPlot, gammaWilsonPlots});

```

It indeed gets the infinite-dilution values right, but overall the fit is incredibly bad!

Try again to determine coefficients, now fitting to complete set of activity-coefficient data.

```

> objective := (L12,L21) -> sum(
  evalf(ln(gamma1Wilson(x1['i'],L12,L21)/gamma1['i'])^2 +
  ln(gamma2Wilson(x1['i'],L12,L21)/gamma2['i'])^2),'i'=1..n);
objective := (L12, L21) ->

$$\sum_{i=1}^n \text{evalf}\left(\ln\left(\frac{\text{gamma1Wilson}(x1_{i}, L12, L21)}{\gamma1_{i}}\right)^2 + \ln\left(\frac{\text{gamma2Wilson}(x1_{i}, L12, L21)}{\gamma2_{i}}\right)^2\right)$$

> objective(112,121);
7.252614302
Minimize the objective function with respect to the lambda's, taking twenty values of each
between zero and 20
> crudeMinimize(objective,20,[0.0001,20],[0.0001,20]);
[2.000090000, 1.000095000, .2421139087]
> l12 := 2.4; l21 := 0.95;
l12 := 2.4
l21 := .95
Now refine the estimates near the rough minimum
> crudeMinimize(objective,10,[1,3],[.5,2]);
[2.400000000, .9500000000, .05387353182]
Here are the best-fit values. Let's try them out.
> l12 := 2.4; l21 := 0.95;
l12 := 2.4
l21 := .95
> gEWilsonPlot := plot(gEWilson(x,l12,l21),x=0..1):
> display({gEWilsonPlot,gEdataPlot});
> gammaWilsonPlots :=
  plot({gamma1Wilson(x,l12,l21),gamma2Wilson(x,l12,l21)},x=0..0
.99):
> display({gamma1dataPlot,gamma2dataPlot,gammaWilsonPlots});
Much better!

```

## Consistency test

```

> gam1inf := gamma1fit(0); gam2inf := gamma2fit(1);
gam1inf := .4769708028
gam2inf := .2868774953
> lnRatio := [seq(ln(gamma1[i]/gamma2[i]),i=1..n)]:
> plotdata(1,n,x1,lnRatio);
> trapezoid := (i0,i1,x,y) ->
  sum(0.5*(y['i'+1]+y['i'])*(x['i'+1]-x['i']),'i'=i0..i1-1);
trapezoid := (i0, i1, x, y) -> 
$$\sum_{i=i0}^{i1-1} (.5 (y_{i+1} + y_i) (x_{i+1} - x_i))$$

> trapezoid(1,n,x1,lnRatio);
-.05366707990
> -trapezoid(1,9,x1,lnRatio)+trapezoid(9,n,x1,lnRatio);
.5074183619

```

Here we compare the adifference in the area above and below to the total area above plus below. Should be less than 0.02 or so.

```

> abs(" ")/";
.1057649544

```

└─ The data don't pass the test!

## ▣ VLE

Here we examine the 2-parameter Margules model as to how well it re-creates the original P-x-y data. We also examine the prediction of the ideal-solution model.

```
> PfitMargules2 := (x,A12,A21) ->
  plsatsat*x*gamma1Margules2(x,A12,A21) +
  p2satsat*(1-x)*gamma2Margules2(x,A12,A21);
PfitMargules2 := (x, A12, A21) →
  plsatsat x gamma1Margules2(x, A12, A21) + p2satsat (1 - x) gamma2Margules2(x, A12, A21)
> yfitMargules2 := (x,A12,A21) ->
  x*gamma1Margules2(x,A12,A21)*plsatsat/PfitMargules2(x,A12,A21);
yfitMargules2 := (x, A12, A21) →  $\frac{x \text{ gamma1Margules2}(x, A12, A21) \text{ plsatsat}}{\text{PfitMargules2}(x, A12, A21)}$ 
> PvsxFit := plot(PfitMargules2(x,a12,a21),x=0..1):
> PvsyFit :=
  plot([yfitMargules2(x,a12,a21),PfitMargules2(x,a12,a21),x=0..1]
  ):
> display({PvsxFit,Pvsx,PvsyFit,Pvsy});
```

## ▣ Ideal solution

```
> PIdeal := x -> x*plsatsat + (1-x)*p2satsat;
yIdeal := x -> x*plsatsat/PIdeal(x);
PIdeal := x → x plsatsat + (1 - x) p2satsat
yIdeal := x →  $\frac{x \text{ plsatsat}}{\text{PIdeal}(x)}$ 
> PvsxIdeal := plot(PIdeal(x),x=0..1):
> PvsyIdeal := plot([yIdeal(x),PIdeal(x),x=0..1]):
> display({PvsxIdeal,Pvsx,PvsyIdeal,Pvsy});
```

## ▣ "Barker's method" applied to 2-constant Margules formula

Here we try Barker's method (discussed in the text). The basic idea is to fit the activity-coefficient model to the P-x data only.

This involves finding the values of the Margules parameters that minimizes the sum-of-square deviation between the model

pressure and the actual data for the pressure.

```
> objective := (A12,A21) -> sum((PfitMargules2(x1['i'],A12,A21)
  - P['i'])^2,'i'=2..n-1);
```

$$\text{objective} := (A12, A21) \rightarrow \sum_{i=2}^{n-1} (\text{PfitMargules2}(x_{I_i}, A12, A21) - P_i)^2$$

```
> print(a12,a21);
```

-0.7403, -1.2487

```
> objective(a12,a21);
```

2.864612930

```
> optimum :=
```

```
  crudeMinimize(objective,20,[0.5*a12,2*a12],[0.5*a21,2*a21]);
```

```
  optimum := [-.7032850000, -1.373570000, .7399279830]
```

```
> a12 := optimum[1]; a21 := optimum[2];
```

a12 := -0.7032850000

a21 := -1.373570000

```

> optimum :=
  crudeMinimize(objective,20,[0.8*a12,1.2*a12],[0.8*a21,1.2*a21
  ]);
                                optimum := [-.7173507000, -1.373570000, .7204838368]
> a12 := optimum[1]; a21 := optimum[2];
                                a12 := -.7173507000
                                a21 := -1.373570000
[ Not a dramatic difference, but visible
> gammaMargules2Plots :=
  plot({gamma1Margules2(x,a12,a21),gamma2Margules2(x,a12,a21)},
  x=0..1):
> display({gamma1dataPlot,gamma2dataPlot,gammaMargules2Plots});
> PvsxFitBarker :=
  plot(PfitMargules2(x,a12,a21),x=0..1,color=green):
> PvsyFitBarker :=
  plot([yfitMargules2(x,a12,a21),PfitMargules2(x,a12,a21),x=0..
  1],color=green):
> display({PvsxFit,Pvsx,PvsyFit,Pvsy,PvsxFitBarker,PvsyFitBarke
  r});

```