

```

In[21]:= experiments = {
    "EYrainbow_glucose",
    "EYrainbow_glucose_largerBF",
    "EYrainbow_rapamycin_1stTry",
    "EYrainbow_rapamycin_CheckBistability",
    "EYrainbow_1nmpp1_1st",
    "EYrainbow_leucine_large",
    "EYrainbowWhi5Up_betaEstrodiol"
};
data = Import[FileNameJoin[{NotebookDirectory[], "fitCellSizeWithOrganelle_normal.csv"}],
    "Data", "HeaderLines" → 1];
epsilon = 10^(-18);
filled = data /. {0. → epsilon};
washed = Select[data, Function[x, NoneTrue[x, # == 0 &]]];

In[ ]:= (* Check data *)
Position[filled, 0.]
Position[washed, 0.]

In[26]:= {}
Out[26]= {}

In[ ]:= Length[filled]
Length[washed]

Out[ ]= 9744

Out[ ]= 6214

In[27]:= fitCellSizeFromOrganelles[dataset_] := Module[
    {expr, a, b, c, alpha, mean, n, nlm, params,
    fittedFunc, fitted, fittedMin, fittedMax, dataMin, dataMax},
    expr = Sum[a[i] × n[i] × mean[i]^alpha[i], {i, 1, 6}] + Sum[
        n[i] × mean[i]^alpha[i] × b[i, j] × n[j] × mean[j]^alpha[j], {i, 1, 6}, {j, i, 6}] + c;
    nlm = NonlinearModelFit[
        dataset[[All, 2 ;;]], expr,
        Join[
            Table[a[i], {i, 1, 6}],
            Flatten[Table[b[i, j], {i, 1, 6}, {j, i, 6}]],
            {c},
            Table[alpha[i], {i, 1, 6}]
        ],
        Join[
            Table[mean[i], {i, 1, 6}],
            Table[n[i], {i, 1, 6}]
        ],
        MaxIterations → ∞

```

```

];
params = nlm["BestFitParameters"];
Print["a[i]: ", Table[a[i], {i, 1, 6}] /. params];
Print["b[i,j]: ", Table[b[i, j], {i, 1, 6}, {j, 1, 6}] /. params // MatrixForm];
Print["c: ", c /. params];
Print["alpha[i]: ", Table[alpha[i], {i, 1, 6}] /. params];
Print["Adjusted R Square: ", nlm["AdjustedRSquared"]];
Print["p-values for parameter z-statistics: ", nlm["ParameterPValues"]];
Print["t-statistics for parameter estimates: ", nlm["ParameterTStatistics"]];
fittedFunc = Function[
  vec,
  expr /. params
  /. Table[n[i] → vec[[i]], {i, 1, 6}]
  /. Table[mean[i] → vec[[i + 6]], {i, 1, 6}]
];
fitted = Map[fittedFunc, dataset[[All, 2 ;; -2]]];
fittedMin = Min[fitted];
fittedMax = Max[fitted];
dataMin = Min[dataset[[All, -1]]];
dataMax = Max[dataset[[All, -1]]];
Print[Show[
  ListPlot[
    Transpose[{dataset[[All, -1]], fitted}],
    PlotTheme → "Scientific"
  ],
  Plot[x, {x, 0, dataMax}]
]];
Print[""]
]

```

```

In[28]:= Print["All Experiments, Null Organelles filled with 0:"];
fitCellSizeFromOrganelles[filled];

```

All Experiments, Null Organelles filled with 0:

a[i]: {0.696729, -9.57666, 0.402288, 0.4979, 1.4033, -0.706302}

b[i,j]:

	0.170765	-0.579225	-0.139695	-0.00518762	0.260611	-0.102986
b\$11540[2, 1]	10.5245	-1.18303	-0.00946476	-0.172667	0.452603	
b\$11540[3, 1]	b\$11540[3, 2]	0.393994	0.00240183	-0.328545	0.154927	
b\$11540[4, 1]	b\$11540[4, 2]	b\$11540[4, 3]	-0.00432734	0.0187626	-0.0257911	
b\$11540[5, 1]	b\$11540[5, 2]	b\$11540[5, 3]	b\$11540[5, 4]	0.191292	-0.22388	
b\$11540[6, 1]	b\$11540[6, 2]	b\$11540[6, 3]	b\$11540[6, 4]	b\$11540[6, 5]	0.0597549	

c: 37.7287

alpha[i]: {0.925117, 0.379822, 0.869212, -0.914228, 0.689248, 1.54461}

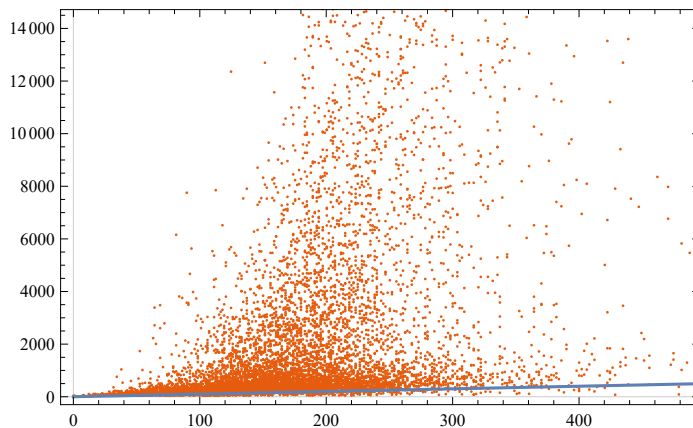
Adjusted R Square: 0.962006

p-values for parameter z-statistics:

{0.0956655, 8.46231×10^{-8} , 0.623104, 1.2917×10^{-8} , 5.00977×10^{-6} , 0.163537, 1.54544×10^{-14} , 1.8314×10^{-11} , 1.77948×10^{-6} , 0.430493, 3.68364×10^{-30} , 0.00146404, 3.90237×10^{-52} , 7.36702×10^{-8} , 0.534402, 0.00104466, 5.87903×10^{-7} , 0.00462498, 0.613966, 1.05604×10^{-9} , 0.000153857, 0.0000151448, 4.55863×10^{-6} , 0.000442365, 6.06702×10^{-88} , 1.00666×10^{-14} , 0.000500276, 7.7456×10^{-38} , 1.85604×10^{-194} , 0., 4.0401×10^{-114} , 2.48635×10^{-52} , 0., 2.45062×10^{-26} }

t-statistics for parameter estimates:

{1.6664, -5.36108, 0.491472, 5.69205, 4.56695, -1.39338, 7.69561, -6.72686, -4.77991, -0.788382, 11.4497, -3.18263, 15.2847, -5.38614, -0.621322, -3.27921, 4.99849, 2.83271, 0.504437, -6.10671, 3.78625, -4.32877, 4.58674, -3.51463, 20.0845, -7.75056, 3.48178, 12.9134, 30.4393, 66.2016, 23.0099, -15.3147, 106.338, 10.6494}



```
In[30]:= Print["All Experiments, Null Organelles Excluded:"];
fitCellSizeFromOrganelles[washed];
```

All Experiments, Null Organelles Excluded:

a[i]: {1.43319, -0.631799, -2.43518, 0.602025, 1.74968, -1.72643}

b[i,j]:

	0.095037	-0.68094	-0.0506076	0.00784741	0.134502	-0.0180395
b\$12022[2, 1]	11.5453	-1.83976	0.102965	-0.485344	0.456931	
b\$12022[3, 1]	b\$12022[3, 2]	0.758316	-0.0382842	-0.376174	0.11432	
b\$12022[4, 1]	b\$12022[4, 2]	b\$12022[4, 3]	-0.0522034	0.109615	-0.0802438	
b\$12022[5, 1]	b\$12022[5, 2]	b\$12022[5, 3]	b\$12022[5, 4]	0.147011	-0.101026	
b\$12022[6, 1]	b\$12022[6, 2]	b\$12022[6, 3]	b\$12022[6, 4]	b\$12022[6, 5]	0.0245043	

c: 41.8575

alpha[i]: {0.97367, 0.378966, 0.810015, -0.203626, 0.727264, 1.81137}

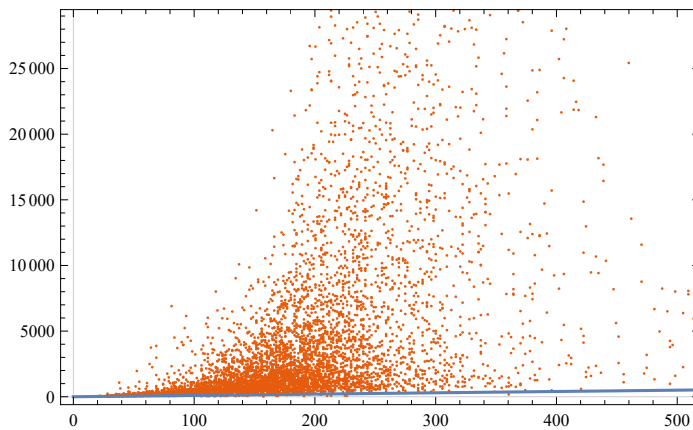
Adjusted R Square: 0.970595

p-values for parameter z-statistics:

{0.0060045, 0.842294, 0.203612, 0.0186789, 8.05145×10^{-7} , 0.0107173, 0.0000359819, 3.40323×10^{-10} , 0.217337, 0.673792, 1.21553×10^{-7} , 0.626614, 2.10451×10^{-18} , 1.11682×10^{-6} , 0.0576047, 4.4877×10^{-11} , 0.000962163, 0.00613598, 0.0644377, 5.73774×10^{-9} , 0.0391787, 1.47696×10^{-12} , 1.01847×10^{-21} , 0.00582111, 2.2024×10^{-44} , 0.00205146, 0.204926, 7.65658×10^{-13} , 5.4706×10^{-118} , 0., 1.80404×10^{-95} , 0.000133278, 0., 1.01369×10^{-10} }

t-statistics for parameter estimates:

{2.74849, -0.198968, -1.27146, 2.3525, 4.93918, -2.55256, 4.1349, -6.28948, -1.23377, 0.42097, 5.29736, -0.486523, 8.77909, -4.87473, 1.89905, -6.59886, 3.30295, 2.74137, -1.84947, -5.83247, 2.06274, -7.09128, 9.61102, -2.75865, 14.0871, -3.08399, 1.26778, 7.18238, 23.601, 42.7854, 21.098, -3.82273, 100.211, 6.4761}



```
In[32]:= Do[
  Print[experiments[[i]], ", Null Organelles filled with 0:"];
  fitCellSizeFromOrganelles[Select[filled, #[[1]] == experiments[[i]] &]];
  {i, 1, Length[experiments]}
];
EYrainbow_glucose, Null Organelles filled with 0:
a[i]: {7.95055×106, -1.87231×106, -1.67851×107, 1.27235×107, -899192., 5.71049×106}
```

$$b[i,j]: \begin{pmatrix} 0.206655 & -2.26605 & -7.95056 \times 10^6 & -0.719441 & -0.330848 & 1.98234 \\ b\$12504[2,1] & 3.20509 & 1.87232 \times 10^6 & 1.43031 & -0.170369 & -0.855271 \\ b\$12504[3,1] & b\$12504[3,2] & 673.245 & -1.27235 \times 10^7 & 899.190 & -5.71051 \times 10^6 \\ b\$12504[4,1] & b\$12504[4,2] & b\$12504[4,3] & -0.0905509 & -0.0507523 & -1.10097 \\ b\$12504[5,1] & b\$12504[5,2] & b\$12504[5,3] & b\$12504[5,4] & 0.145153 & -0.237671 \\ b\$12504[6,1] & b\$12504[6,2] & b\$12504[6,3] & b\$12504[6,4] & b\$12504[6,5] & 0.279037 \end{pmatrix}$$

c: 1.6112×10^7

alpha[i]: {0.711231, 0.472112, -1.01925×10^{-6} , 0.528598, 0.857688, 0.518671}

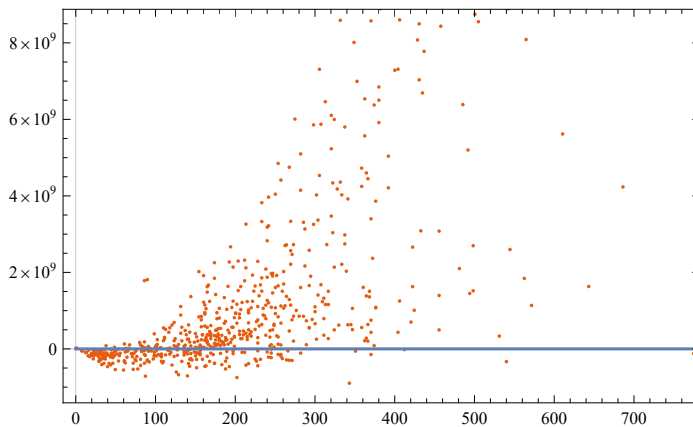
Adjusted R Square: 0.957991

p-values for parameter z-statistics:

{0., 0., 0., 0., 0., 0., 0.558229, 0.0000385444, 0., 0.0349717, 0.0286449,
0.00161899, 0.0150112, 0., 0.000040135, 0.150444, 0.0277493, 0., 0., 0., 0., 0.482971,
0.569478, 0.00805373, 0.000044358, 0.0599673, 0.221337, 0., 6.81699×10^{-23} ,
 1.44105×10^{-40} , 2.46376×10^{-16} , 1.09666×10^{-6} , 1.96383×10^{-186} , 0.0117437}

t-statistics for parameter estimates:

{ 3.51057×10^6 , -737.628., -4.00951×10^6 , 5.29553×10^6 , -1.6491×10^6 , 3.41428×10^6 , 0.585788,
-4.14572, -3.51058×10^6 , -2.11328, -2.19341, 3.16629, 2.43884, 737.630., 4.13619, -1.43973,
-2.20596, 160.820., -5.29556×10^6 , 1.64914×10^6 , -3.41429×10^6 , -0.701951, -0.569121, -2.65832,
4.11253, -1.88445, 1.2242, 3.8487×10^6 , 10.2493, 14.3474, -8.42551, 4.92204, 42.4541, 2.52717}



EYrainbow_glucose_largerBF, Null Organelles filled with 0:

a[i]: {-4.81783, 16.4708, -162.458, -4.4573, 7.03307, 8.63351}

$$b[i,j]: \begin{pmatrix} 0.407012 & -4.54289 & 1.91442 & -0.0795996 & 0.095331 & 0.706776 \\ b\$12662[2,1] & 17.2503 & -21.7685 & 2.07376 & -0.323612 & 0.194778 \\ b\$12662[3,1] & b\$12662[3,2] & 88.1717 & 1.82106 & -5.43396 & -2.26305 \\ b\$12662[4,1] & b\$12662[4,2] & b\$12662[4,3] & 0.00428663 & -0.594038 & 0.724965 \\ b\$12662[5,1] & b\$12662[5,2] & b\$12662[5,3] & b\$12662[5,4] & 0.359781 & -0.385994 \\ b\$12662[6,1] & b\$12662[6,2] & b\$12662[6,3] & b\$12662[6,4] & b\$12662[6,5] & 0.0978354 \end{pmatrix}$$

c: 59.8892

alpha[i]: {0.619894, 0.329772, 0.279838, 1.88814, 0.729227, 0.494416}

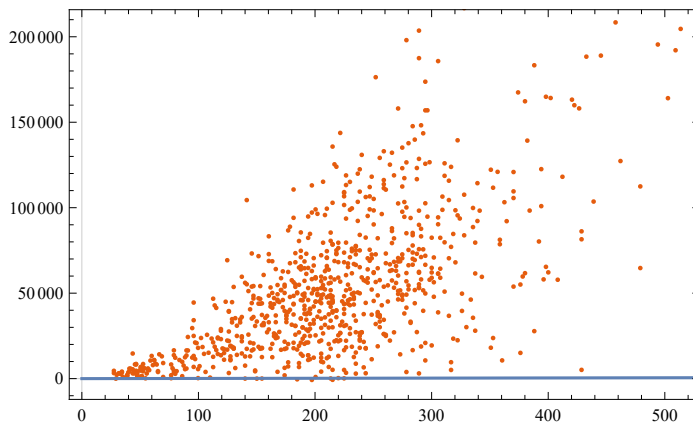
Adjusted R Square: 0.96968

p-values for parameter z-statistics:

```
{0.655467, 0.600074, 0.665985, 0.513876, 0.319762, 0.168583, 0.283156, 1.2888×10-6,
0.757159, 0.850959, 0.665257, 0.0509041, 0.000417492, 0.368162, 0.00354378,
0.269127, 0.669582, 0.62403, 0.681485, 0.335958, 0.536978, 0.981929,
0.000668084, 0.0111833, 9.15186×10-18, 0.0166275, 0.543321, 0.757275,
1.2395×10-15, 7.60049×10-27, 0.0994903, 1.77737×10-8, 4.75916×10-234, 0.00141722}
```

t-statistics for parameter estimates:

```
{-0.446342, 0.524497, -0.43182, -0.653096, 0.995545, 1.37798, 1.07395, -4.87747,
0.309316, -0.187953, 0.432822, 1.95513, 3.54318, -0.900414, 2.92455, -1.10582, 0.426875,
0.49033, 0.410583, -0.962742, -0.617647, 0.0226575, -3.41533, 2.54263, 8.78171,
-2.39973, 0.608051, 0.309163, 8.16034, 11.1148, 1.64918, 5.68858, 46.6933, 3.20194}
```



EYrainbow_rapamycin_1stTry, Null Organelles filled with 0:

a[i]: {3.42279, -13.3567, 0.685418, -9.61768, 7.04366, -2.07536}

	0.313322	-0.495174	-0.018953	-0.414813	0.328839	-0.360455
b\$12800[2, 1]	9.10798	-0.321376	1.47292	-1.18016	0.130376	
b\$12800[3, 1]	b\$12800[3, 2]	0.0767022	-0.0217186	-0.133651	-0.0835466	
b\$12800[4, 1]	b\$12800[4, 2]	b\$12800[4, 3]	0.379327	0.0985121	0.189071	
b\$12800[5, 1]	b\$12800[5, 2]	b\$12800[5, 3]	b\$12800[5, 4]	0.11439	0.0194904	
b\$12800[6, 1]	b\$12800[6, 2]	b\$12800[6, 3]	b\$12800[6, 4]	b\$12800[6, 5]	0.216252	

c: 62.5259

alpha[i]: {0.37867, 0.403837, 1.04573, 2.64262, 0.303223, 0.435492}

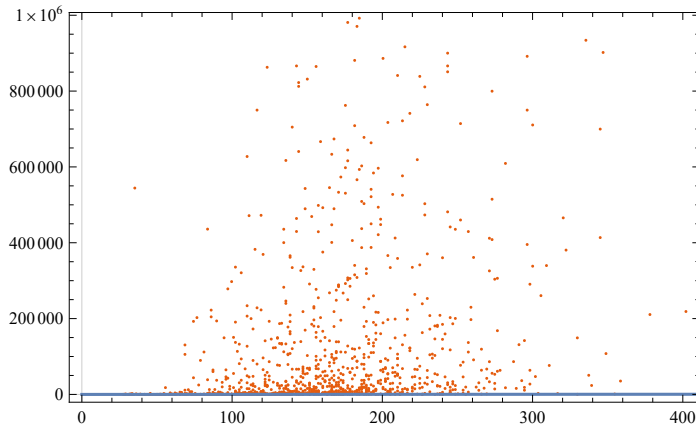
Adjusted R Square: 0.953756

p-values for parameter z-statistics:

```
{0.0342084, 0.00430735, 0.34639, 0.00180234, 1.17166×10-6, 0.0527066, 0.110936, 0.218935, 0.762829,
0.23231, 0.100213, 0.0053468, 0.0000746227, 0.21696, 0.0174269, 0.000169527, 0.469685, 0.526541,
0.786469, 0.207605, 0.261278, 0.169067, 0.7128, 0.256484, 0.293293, 0.829232, 0.000769661,
2.62782×10-34, 0.000779044, 4.09033×10-43, 3.98661×10-9, 7.45811×10-7, 2.57607×10-6, 0.00373028}
```

t-statistics for parameter estimates:

```
{2.11994, -2.85999, 0.941971, -3.12776, 4.88453, -1.93917, 1.59513, -1.22999,
-0.301835, -1.19501, 1.64504, -2.79027, 3.97437, -1.23528, 2.38076, -3.77204,
0.723212, 0.633473, -0.270957, -1.26083, -1.12389, 1.37601, 0.368183, 1.13526, 1.05137,
0.215732, 3.37182, 12.5918, 3.36845, 14.3102, 5.92715, 4.97451, 4.72394, 2.90561}
```



EYrainbow_rapamycin_CheckBistability, Null Organelles filled with 0:

a[i]: {-3.74637, -22.4408, 1.73533, 4.9535, 0.490163, -0.380536}

b[i,j]:

0.219122	-0.0306399	-0.232835	-0.263235	0.37083	-0.274052
b\$12938[2, 1]	15.2545	-2.08231	0.34816	0.317638	-0.086636
b\$12938[3, 1]	b\$12938[3, 2]	1.45838	-1.50555	-0.886865	0.307814
b\$12938[4, 1]	b\$12938[4, 2]	b\$12938[4, 3]	1.3352	0.306796	0.284691
b\$12938[5, 1]	b\$12938[5, 2]	b\$12938[5, 3]	b\$12938[5, 4]	0.226377	-0.22029
b\$12938[6, 1]	b\$12938[6, 2]	b\$12938[6, 3]	b\$12938[6, 4]	b\$12938[6, 5]	0.0858692

c: 34.1267

alpha[i]: {1.01523, 0.32123, 0.781379, 0.629678, 0.832371, 0.75963}

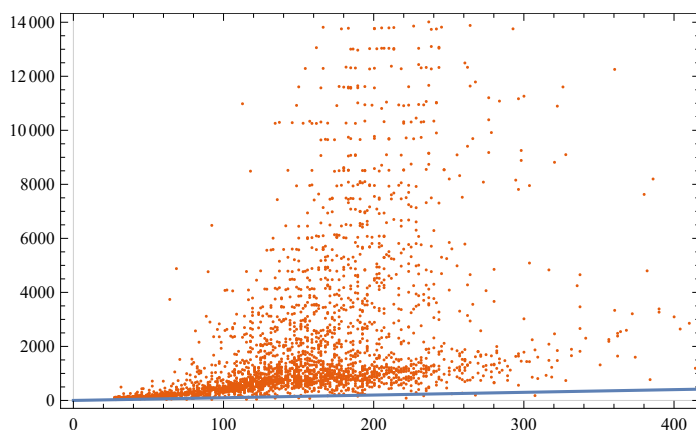
Adjusted R Square: 0.971839

p-values for parameter z-statistics:

{5.13048×10⁻⁷, 2.13525×10⁻⁸, 0.521037, 0.0208712, 0.507063, 0.601608, 2.37029×10⁻⁹, 0.842149, 0.00870986, 0.0241644, 2.17033×10⁻¹⁶, 1.4093×10⁻⁹, 2.54424×10⁻¹², 0.000139829, 0.371839, 0.0215657, 0.539243, 0.0282786, 0.000448397, 2.0767×10⁻⁶, 0.00119048, 0.00250105, 0.00314945, 0.00648853, 3.72076×10⁻¹⁶, 3.36409×10⁻⁸, 0.000356346, 6.13713×10⁻¹⁰, 7.56557×10⁻⁹⁶, 1.45827×10⁻⁷⁹, 4.84154×10⁻⁵⁴, 1.46333×10⁻⁷, 0., 2.83143×10⁻¹⁴}

t-statistics for parameter estimates:

{-5.03195, -5.61525, 0.641821, 2.31148, 0.663495, -0.522143, 5.98776, -0.199162, -2.6249, -2.25561, 8.2576, -6.07276, 7.02904, -3.81321, 0.893163, 2.29909, -0.614026, 2.19442, -3.51358, -4.75479, 3.24415, 3.02568, 2.95507, 2.72385, 8.19158, -5.53533, 3.57441, 6.20642, 21.5157, 19.4425, 15.7823, 5.26922, 53.9007, 7.64187}



EYrainbow_1nmppl_1st, Null Organelles filled with 0:

$a[i]: \{4.1015, -7.49712, 2.17978, 1.02, 3.65027, -4.75357\}$

$b[i,j]:$

	0.190813	-0.347852	-0.0341533	0.19114	-0.0894577	-0.234675
$b\$13076[2, 1]$	10.3124	-0.784643	1.09945	-0.182763	0.124003	
$b\$13076[3, 1]$	$b\$13076[3, 2]$	0.116329	-0.20437	-0.245506	0.010035	
$b\$13076[4, 1]$	$b\$13076[4, 2]$	$b\$13076[4, 3]$	0.0876259	0.125328	0.234293	
$b\$13076[5, 1]$	$b\$13076[5, 2]$	$b\$13076[5, 3]$	$b\$13076[5, 4]$	0.212202	0.0369924	
$b\$13076[6, 1]$	$b\$13076[6, 2]$	$b\$13076[6, 3]$	$b\$13076[6, 4]$	$b\$13076[6, 5]$	0.11108	

$c: -8.8883$

$\alpha[i]: \{1.17032, 0.404532, 1.0571, 0.0872777, 0.715148, 0.930041\}$

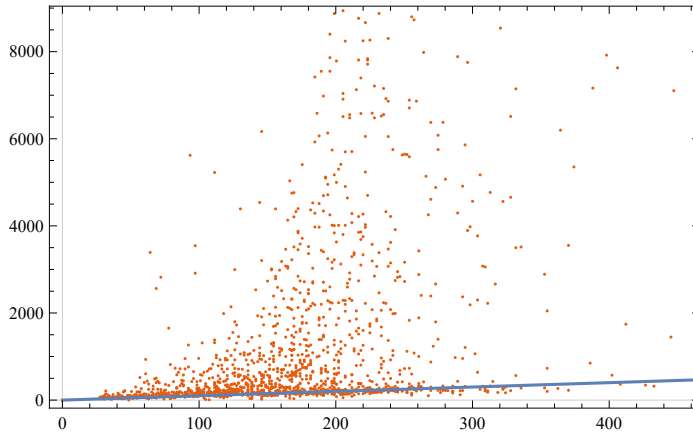
Adjusted R Square: 0.969918

p-values for parameter z-statistics:

$\{0.00493782, 0.0561456, 3.5129 \times 10^{-7}, 0.379877, 0.00054923, 0.000325854, 0.0477486,$
 $0.207459, 0.400098, 0.108, 0.334648, 0.0616406, 8.27744 \times 10^{-9}, 0.00304734,$
 $0.000305166, 0.357422, 0.625858, 0.0930838, 0.00753064, 0.000210796, 0.831232,$
 $0.164733, 0.16541, 0.0434917, 5.04474 \times 10^{-8}, 0.687586, 0.306654, 0.15955,$
 $6.26302 \times 10^{-16}, 1.58833 \times 10^{-94}, 4.13454 \times 10^{-52}, 0.534286, 2.68973 \times 10^{-157}, 0.000791547\}$

t-statistics for parameter estimates:

$\{2.81496, -1.91129, 5.11506, 0.878365, 3.46249, -3.60163, 1.98112, -1.26109,$
 $-0.841669, 1.60814, -0.965083, -1.87018, 5.79354, -2.96744, 3.61878, -0.920554,$
 $0.487659, 1.68035, -2.67581, -3.71423, 0.213157, 1.38996, 1.38774, 2.02055, 5.47588,$
 $0.402205, 1.02259, -1.40724, 8.16833, 22.0748, 15.7546, 0.62161, 29.9908, 3.36215\}$



EYrainbow_leucine_large, Null Organelles filled with 0:

a[i]: {0.0049408, 0.937427, 1.88228, -0.49819, -0.0698077, 0.896746}

b[i,j]:

3.45922×10^{-8}	0.0000105572	-0.0000362882	-0.000274834	0.0000276168	-0.000499061
b\$13214[2, 1]	0.440096	-0.0983444	0.07909	0.00596252	-0.00415221
b\$13214[3, 1]	b\$13214[3, 2]	0.0209771	-0.0313891	-0.00131623	-0.0342072
b\$13214[4, 1]	b\$13214[4, 2]	b\$13214[4, 3]	0.0292557	0.0000688139	0.0686443
b\$13214[5, 1]	b\$13214[5, 2]	b\$13214[5, 3]	b\$13214[5, 4]	0.0000466083	0.00332761
b\$13214[6, 1]	b\$13214[6, 2]	b\$13214[6, 3]	b\$13214[6, 4]	b\$13214[6, 5]	0.0580782

c: 15.2346

alpha[i]: {4.83104, 0.659347, 1.15183, 0.418643, 1.61619, 0.305701}

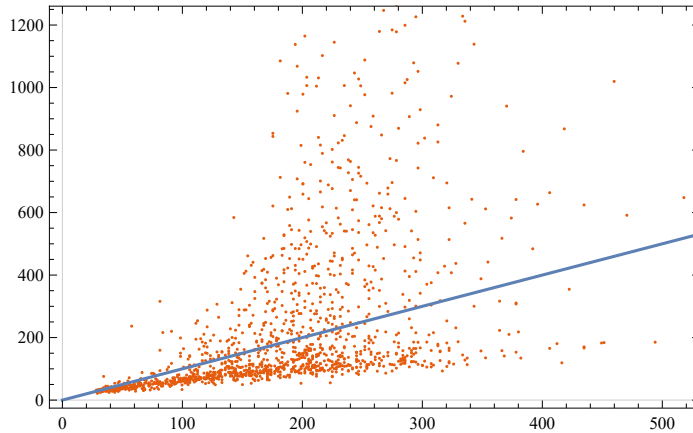
Adjusted R Square: 0.980936

p-values for parameter z-statistics:

{0.650029, 0.616572, 0.00766671, 0.359438, 0.526426, 0.213347, 0.843787, 0.983403, 0.786603,
0.626136, 0.746296, 0.659109, 0.111455, 0.399697, 0.0351601, 0.520874, 0.916808, 0.679908,
0.370178, 0.584709, 0.364169, 0.00260943, 0.982838, 0.0130355, 0.740785, 0.461298, 0.0293473,
0.000275306, 0.000137445, 7.48556×10^{-22} , 1.09958×10^{-6} , 0.110603, 0.0000318801, 0.00137052}

t-statistics for parameter estimates:

{0.453815, 0.500821, 2.67009, -0.916723, -0.633622, 1.24495, 0.197087, 0.0208061, -0.270773,
-0.487275, 0.323588, -0.441234, 1.59264, -0.842407, 2.10846, 0.642153, -0.104473,
0.41267, -0.896414, -0.546641, -0.907731, 3.01551, 0.0215145, 2.48583, 0.330876,
0.736905, 2.18092, 3.64631, 3.82281, 9.76115, 4.89349, 1.59645, 4.17258, 3.20692}



EYrainbowWhi5Up_betaEstradiol, Null Organelles filled with 0:

a[i]: {10.2175, 1.17026, -120.585, 7.73935, 9.92105, -7.07683}

b[i,j]:

	0.239493	-1.59017	-2.05651	0.0418467	0.229477	-0.341809
b\$13352[2, 1]		8.90769	-1.44567	1.36135	-1.29201	0.0498444
b\$13352[3, 1]	b\$13352[3, 2]		35.9827	-2.94319	-4.29195	-0.122284
b\$13352[4, 1]	b\$13352[4, 2]	b\$13352[4, 3]		0.19809	-0.119025	-0.372668
b\$13352[5, 1]	b\$13352[5, 2]	b\$13352[5, 3]	b\$13352[5, 4]		0.465209	-0.0700564
b\$13352[6, 1]	b\$13352[6, 2]	b\$13352[6, 3]	b\$13352[6, 4]	b\$13352[6, 5]		0.813159

c: 91.2457

alpha[i]: {1.04828, 0.412252, 0.419638, 1.44656, 0.704655, 1.33722}

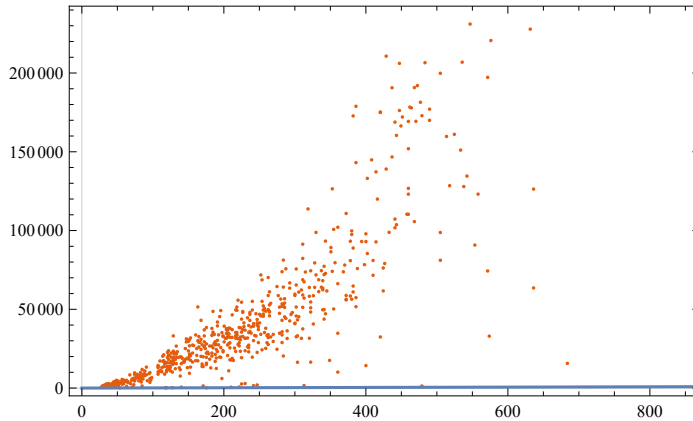
Adjusted R Square: 0.981235

p-values for parameter z-statistics:

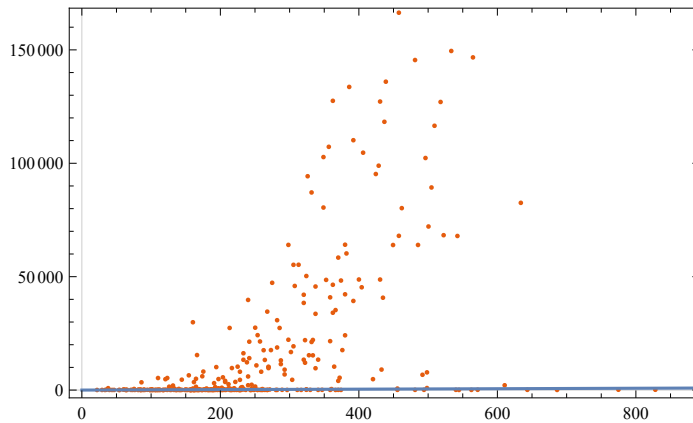
{0.00354431, 0.878666, 0.416446, 0.193492, 0.0113249, 0.115137, 0.00145578,
 1.145×10^{-7} , 0.153766, 0.835776, 0.0747082, 0.0530184, 0.0000179972, 0.568837,
 0.00526238, 0.000317945, 0.910876, 0.417664, 0.240857, 0.0735164, 0.932812,
 0.322244, 0.635695, 0.241769, 0.00371187, 0.771323, 0.000812342, 0.487042,
 1.60005×10^{-48} , 8.74393×10^{-87} , 0.0002131, 9.79986×10^{-6} , 7.08386×10^{-101} , 1.76366×10^{-6} }

t-statistics for parameter estimates:

{2.92737, 0.152724, -0.813149, 1.30174, 2.54014, -1.57774, 3.19778, -5.36525,
 -1.42811, 0.207387, 1.78527, -1.9385, 4.32232, -0.570071, 2.80049, -3.62069, 0.111979,
 0.811024, -1.17398, -1.79266, -0.0843422, 0.990651, -0.473961, -1.1717, 2.91278,
 -0.290772, 3.3652, 0.69544, 16.0136, 23.3296, 3.72515, 4.45863, 25.9391, 4.82532}



```
In[33]:= Do[
  Print[experiments[[i]], ", Null Organelles Excluded:"];
  fitCellSizeFromOrganelles[Select[washed, #[[1] == experiments[[i]] &]];
  {i, 1, Length[experiments]}
]
EYrainbow_glucose, Null Organelles Excluded:
a[i]: {5.47487, -27.622, 0.48096, -3.13462, 2.88081×10-6, 3.85831}
b[i,j]: {
  0.445775      -2.35577      -0.0198103      0.322307      -1.78057×10-7      1.03061
  b$13819[2, 1]  10.1805      -0.069881      1.98776      4.42265×10-6      -0.274067
  b$13819[3, 1]  b$13819[3, 2]  0.000454131      -0.00992291      -2.22761×10-7      -0.000682103
  b$13819[4, 1]  b$13819[4, 2]  b$13819[4, 3]      0.141132      -2.35799×10-7      -1.00324
  b$13819[5, 1]  b$13819[5, 2]  b$13819[5, 3]  b$13819[5, 4]  6.91581×10-11  5.90736×10-6
  b$13819[6, 1]  b$13819[6, 2]  b$13819[6, 3]  b$13819[6, 4]  b$13819[6, 5]      0.0202411
}
c: 45.0752
alpha[i]: {0.7555, 0.394194, 1.70892, 0.32841, 3.04428, 0.557125}
Adjusted R Square: 0.97295
p-values for parameter z-statistics:
{0.211749, 0.0611537, 0.221599, 0.284506, 0.954572, 0.37855, 0.191762, 0.00288126, 0.461784,
 0.415901, 0.965293, 0.181418, 0.0271451, 0.429755, 0.00747604, 0.731161, 0.641747, 0.696287,
 0.544998, 0.76166, 0.938985, 0.390977, 0.941976, 0.114978, 0.875476, 0.758778, 0.939913,
 0.0913981, 2.49285×10-9, 4.6641×10-26, 1.2273×10-10, 0.061419, 8.01698×10-6, 0.189896}
t-statistics for parameter estimates:
{1.25115, -1.87866, 1.22455, -1.07196, 0.0570091, 0.881732, 1.30801, -3.00212, -0.736754,
 0.814558, -0.0435451, 1.33917, 2.21911, -0.790564, 2.69113, 0.343866, -0.465678,
 0.390671, -0.605887, -0.303547, -0.0766033, 0.858953, -0.072841, -1.58029, 0.156825,
 0.307335, 0.075435, 1.6929, 6.12739, 11.5041, 6.64402, 1.87673, 4.53502, 1.31354}
```



EYrainbow_glucose_largerBF, Null Organelles Excluded:

$a[i]: \{-45.0484, 33.9864, -570.005, -14.5716, 24.6618, 145.37\}$

$b[i,j]:$

	-0.436723	-5.08567	30.5792	-0.103099	0.271425	7.68666
b\$13956[2, 1]	32.2023	-47.673	4.07912	-1.04797	-20.3763	
b\$13956[3, 1]	b\$13956[3, 2]	360.302	9.15637	-20.7306	-140.159	
b\$13956[4, 1]	b\$13956[4, 2]	b\$13956[4, 3]	-0.0321034	-0.379627	0.808794	
b\$13956[5, 1]	b\$13956[5, 2]	b\$13956[5, 3]	b\$13956[5, 4]	0.319971	3.81128	
b\$13956[6, 1]	b\$13956[6, 2]	b\$13956[6, 3]	b\$13956[6, 4]	b\$13956[6, 5]	-3.15506	

$c: 179.862$

$\alpha[i]: \{0.377443, 0.282921, 0.177476, 1.93791, 0.768022, 3.71759\}$

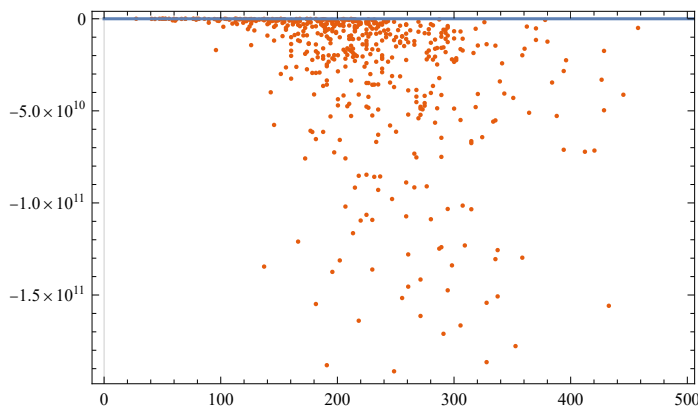
Adjusted R Square: 0.974838

p-values for parameter z-statistics:

$\{0.478609, 0.756873, 0.801825, 0.51458, 0.523898, 0.722741, 0.285719, 0.00909544, 0.589503, 0.811379, 0.200739, 0.60553, 0.119859, 0.615584, 0.0235073, 0.0764494, 0.59506, 0.768026, 0.63639, 0.555947, 0.697689, 0.874972, 0.0200263, 0.86292, 5.00346 \times 10^{-18}, 0.634834, 0.920422, 0.860223, 0.0010816, 1.0835 \times 10^{-7}, 0.398792, 0.000441774, 4.68247 \times 10^{-205}, 0.0563117\}$

t-statistics for parameter estimates:

$\{-0.709021, 0.309743, -0.251106, -0.65214, 0.637751, 0.354978, -1.06859, -2.61763, 0.539869, -0.238761, 1.28097, 0.516758, 1.55776, -0.502401, 2.27135, -1.77497, -0.531824, 0.295103, 0.473012, -0.589229, -0.388644, -0.157418, -2.33258, 0.172738, 8.95724, 0.475195, -0.0999479, 0.176173, 3.28565, 5.38272, 0.84443, 3.53498, 49.2848, 1.91261\}$



EYrainbow_rapamycin_1stTry, Null Organelles Excluded:

a[i]: {4.73974, -19.7446, -3.30983, -4.34326, 10.761, 0.257584}

b[i,j]:

	0.348908	-1.13291	-0.0161668	-0.525506	0.587515	-0.481935
b\$14094[2, 1]	18.5761	-2.39918	2.36921	-1.7913	-0.0635328	
b\$14094[3, 1]	b\$14094[3, 2]	0.576221	-0.0925034	-0.241828	-0.146956	
b\$14094[4, 1]	b\$14094[4, 2]	b\$14094[4, 3]	0.138856	-0.0288575	-0.0377778	
b\$14094[5, 1]	b\$14094[5, 2]	b\$14094[5, 3]	b\$14094[5, 4]	-0.0345846	-0.0538965	
b\$14094[6, 1]	b\$14094[6, 2]	b\$14094[6, 3]	b\$14094[6, 4]	b\$14094[6, 5]	0.271677	

c: 70.2659

alpha[i]: {0.315754, 0.349702, 0.847956, 1.00952, 0.0681449, 0.729603}

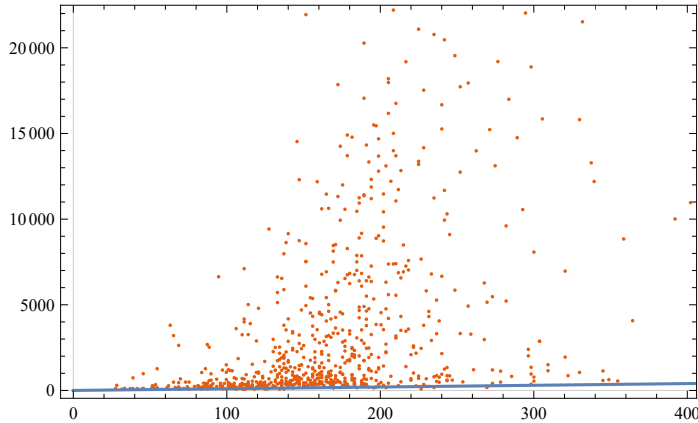
Adjusted R Square: 0.9627

p-values for parameter z-statistics:

{0.0775723, 0.222994, 0.541578, 0.0582602, 1.91954×10^{-8} , 0.881894, 0.122474, 0.169235, 0.938343, 0.0321505, 0.00814444, 0.00640588, 0.0185254, 0.126015, 0.00307349, 0.00307729, 0.87877, 0.408069, 0.538065, 0.237553, 0.321317, 0.292335, 0.865514, 0.771512, 0.781321, 0.637003, 0.00467298, 0.000063441, 0.00881426, 1.13844×10^{-17} , 2.51326×10^{-10} , 0.00126408, 0.553603, 0.000741283}

t-statistics for parameter estimates:

{1.76733, -1.2196, -0.610703, -1.89658, 5.67963, 0.148618, 1.54619, -1.37597, -0.0773786, -2.14647, 2.65298, -2.7338, 2.36001, -1.53168, 2.96979, -2.9694, -0.152579, 0.827753, -0.616021, -1.18205, -0.992399, 1.05374, -0.169416, -0.290501, -0.277696, -0.472083, 2.83712, 4.02202, 2.62594, 8.77161, 6.41218, 3.23609, 0.592631, 3.38758}



EYrainbow_rapamycin_CheckBistability, Null Organelles Excluded:

a[i]: {-0.23803, 2.3262, -4.56638, 2.45662, 3.33995, -0.438266}

b[i,j]:

	0.154752	-0.610254	-0.166578	-0.414718	0.278267	-0.194603
b\$14232[2, 1]	9.01304	-1.895	1.34059	-0.170338	-0.0102316	
b\$14232[3, 1]	b\$14232[3, 2]	1.25754	-1.27152	-0.548598	0.185579	
b\$14232[4, 1]	b\$14232[4, 2]	b\$14232[4, 3]	1.68252	0.232938	0.321991	
b\$14232[5, 1]	b\$14232[5, 2]	b\$14232[5, 3]	b\$14232[5, 4]	0.113859	-0.144862	
b\$14232[6, 1]	b\$14232[6, 2]	b\$14232[6, 3]	b\$14232[6, 4]	b\$14232[6, 5]	0.050741	

c: 18.0912

```
alpha[i]: {1.09905, 0.390899, 0.796849, 0.857423, 0.811011, 0.803621}
```

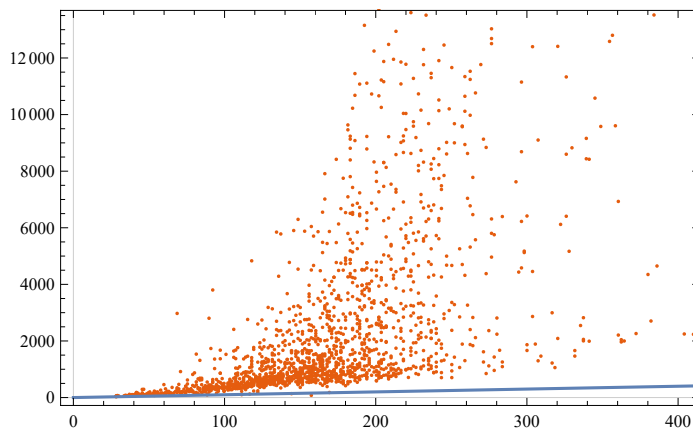
```
Adjusted R Square: 0.984654
```

```
p-values for parameter z-statistics:
```

```
{0.734982, 0.640192, 0.221408, 0.393432, 6.31154×10-8, 0.532498, 1.22048×10-6,  
0.000205366, 0.0255628, 0.00682979, 1.55836×10-10, 1.63753×10-6, 0.00241176, 0.000809832,  
0.0201223, 0.254784, 0.942206, 0.0186756, 0.00142239, 1.49135×10-7, 0.00920578,  
0.00571894, 0.0945132, 0.0150476, 0.0000895055, 0.000383897, 0.0194581, 0.0942608,  
5.94301×10-93, 5.3714×10-33, 1.50846×10-62, 2.73864×10-9, 6.2969×10-196, 2.02219×10-9}
```

```
t-statistics for parameter estimates:
```

```
{-0.338558, 0.467509, -1.22321, 0.853614, 5.43244, -0.624317, 4.86878, -3.71992,  
-2.23464, -2.70814, 6.43682, -4.8096, 3.03848, -3.35507, 2.32612, -1.13917,  
-0.0725072, 2.35407, -3.195, -5.27445, 2.60712, 2.76678, 1.67293, 2.43355, 3.92627,  
-3.5576, 2.33872, 1.67422, 21.694, 12.2081, 17.3576, 5.97654, 33.9515, 6.02675}
```



```
EYrainbow_1nmpp1_1st, Null Organelles Excluded:
```

```
a[i]: {3.50591, 0.897568, 1.61745, 2.68903, 2.30252, -3.56969}
```

```
b[i,j]: {
  {0.00505295, -0.433991, -0.122273, 0.23234, 0.0869233, -0.288756},
  {b$14370[2, 1], 8.59915, -1.41511, 0.710117, 0.11164, -0.336883},
  {b$14370[3, 1], b$14370[3, 2], 0.428631, -0.621625, -0.320523, 0.216963},
  {b$14370[4, 1], b$14370[4, 2], b$14370[4, 3], 0.167718, 0.260291, 0.351818},
  {b$14370[5, 1], b$14370[5, 2], b$14370[5, 3], b$14370[5, 4], 0.073592, -0.117671},
  {b$14370[6, 1], b$14370[6, 2], b$14370[6, 3], b$14370[6, 4], b$14370[6, 5], 0.111619}
}
```

```
c: -6.72342
```

```
alpha[i]: {1.25941, 0.422415, 0.934929, 0.208093, 0.94193, 0.937568}
```

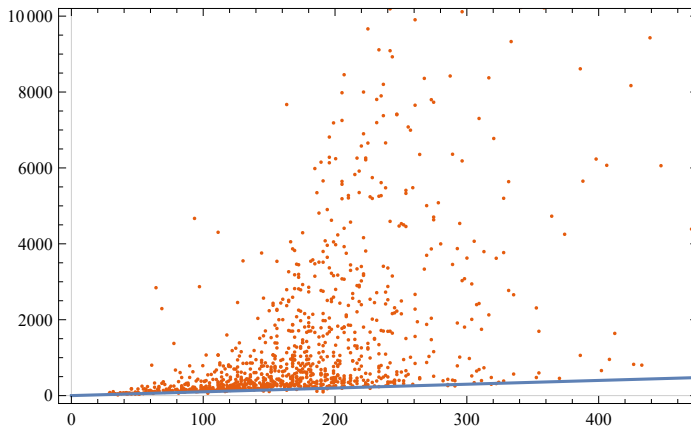
```
Adjusted R Square: 0.976039
```

```
p-values for parameter z-statistics:
```

```
{0.0228285, 0.883692, 0.343123, 0.138008, 0.00189664, 0.0277844, 0.952776, 0.186502,  
0.20622, 0.0916782, 0.152679, 0.0513746, 0.00457972, 0.00597166, 0.0484627,  
0.383931, 0.312617, 0.0848364, 0.00437058, 0.000263823, 0.0769743, 0.080444,  
0.000884034, 0.0162084, 0.00456168, 0.0860328, 0.324016, 0.387998, 2.38466×10-9,  
2.00457×10-35, 1.74046×10-45, 0.0930698, 2.58072×10-104, 0.000854772}
```

t-statistics for parameter estimates:

```
{2.27943, 0.146324, 0.948413, 1.48431, 3.11334, -2.20319, 0.0592347, -1.32181,
-1.26476, 1.68806, 1.4311, -1.95042, 2.84092, -2.75459, 1.97544, 0.871017, -1.0102,
1.72481, -2.8559, -3.66012, 1.77015, 1.7497, 3.334, 2.40783, 2.84219, -1.71822,
0.986673, -0.86359, 6.01823, 12.8638, 14.8146, 1.68085, 24.181, 3.34345}
```



EYrainbow_leucine_large, Null Organelles Excluded:

a[i]: {0.0563306, 2.04486, 1.15134, 1.65373, 0.244904, 0.814932}

	-0.0657253	-0.261834	0.010063	-0.0555318	0.24793	-0.156858
b\$14508[2, 1]	0.929668	-0.0553148	0.0494852	-0.163582	-0.0452303	
b\$14508[3, 1]	b\$14508[3, 2]	0.0080651	-0.00809441	-0.0222354	-0.0125062	
b\$14508[4, 1]	b\$14508[4, 2]	b\$14508[4, 3]	-0.00480908	-0.0798616	0.0784637	
b\$14508[5, 1]	b\$14508[5, 2]	b\$14508[5, 3]	b\$14508[5, 4]	0.14583	-0.139182	
b\$14508[6, 1]	b\$14508[6, 2]	b\$14508[6, 3]	b\$14508[6, 4]	b\$14508[6, 5]	0.132638	

c: 8.73153

alpha[i]: {-0.644591, 0.594088, 1.25099, -0.823046, 0.545933, 0.394545}

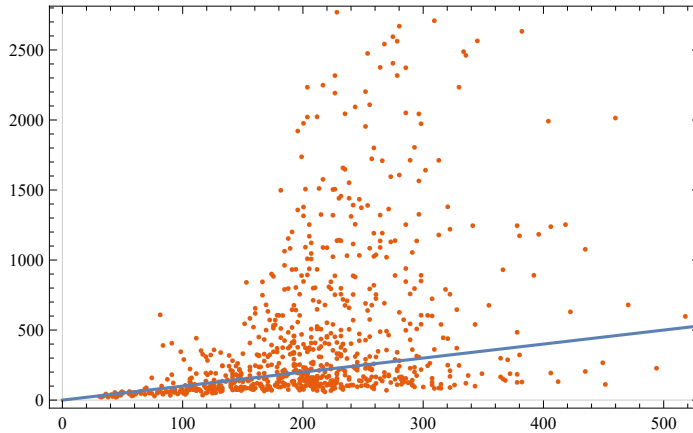
Adjusted R Square: 0.983538

p-values for parameter z-statistics:

```
{0.96106, 0.452549, 0.181731, 0.0117347, 0.847813, 0.362646, 0.286113, 0.0835603, 0.661097,
0.149357, 0.0160564, 0.0648707, 0.163819, 0.600998, 0.307907, 0.155351, 0.530069, 0.802593,
0.622129, 0.600027, 0.610315, 0.682608, 0.059487, 0.0134385, 0.0041345, 0.0718114, 0.00907382,
0.362869, 0.000984223, 3.19194×10-15, 0.00154247, 2.01986×10-8, 1.17755×10-9, 0.000600029}
```

t-statistics for parameter estimates:

```
{0.0488385, 0.751528, 1.33657, 2.52565, 0.19197, 0.91084, -1.06738, -1.73249, 0.438557,
-1.44315, 2.4126, -1.84862, 1.39358, -0.523167, 1.02026, -1.42218, -0.628165,
0.250073, -0.493021, -0.524564, -0.509818, -0.409052, -1.88718, 2.47727, 2.87575,
-1.80263, 2.61545, 0.910415, -3.30688, 8.03648, 3.17719, -5.66597, 6.15403, 3.44493}
```



EYrainbowWhi5Up_betaEstradiol, Null Organelles Excluded:

a[i]: {4.54891, 20.3794, -296.475, 6.26267, 14.5556, -16.4669}

b[i,j]:

0.146396	-1.10991	-0.669914	0.119984	0.168517	-0.401359
b\$14646[2, 1]	6.04524	-6.56303	1.24744	-0.954179	0.364823
b\$14646[3, 1]	b\$14646[3, 2]	91.5896	-2.24536	-7.47329	3.7148
b\$14646[4, 1]	b\$14646[4, 2]	b\$14646[4, 3]	0.117995	-0.420101	-0.153807
b\$14646[5, 1]	b\$14646[5, 2]	b\$14646[5, 3]	b\$14646[5, 4]	0.457192	-0.178164
b\$14646[6, 1]	b\$14646[6, 2]	b\$14646[6, 3]	b\$14646[6, 4]	b\$14646[6, 5]	0.694052

c: 239.26

alpha[i]: {1.18138, 0.438105, 0.333174, 1.87722, 0.741403, 1.51238}

Adjusted R Square: 0.983706

p-values for parameter z-statistics:

{0.281877, 0.150095, 0.582323, 0.439661, 0.0819993, 0.0329787, 0.0132297, 0.0000169754, 0.700281, 0.527166, 0.138474, 0.0146168, 0.00226201, 0.418276, 0.0105491, 0.00282697, 0.377801, 0.603376, 0.547236, 0.24661, 0.379254, 0.500916, 0.129496, 0.677518, 0.00175066, 0.432986, 0.0105446, 0.569403, 1.0339×10^{-40} , 6.78131×10^{-59} , 0.0335698, 2.87383×10^{-6} , 7.74359×10^{-122} , 0.000019766}

t-statistics for parameter estimates:

{1.07714, 1.44116, -0.550314, 0.773315, 1.74232, -2.13761, 2.4854, -4.33889, -0.385138, 0.632732, 1.4836, -2.44931, 3.06754, -0.810012, 2.56588, -2.99915, 0.882653, 0.519845, -0.602271, -1.15983, 0.879965, 0.673483, -1.51832, -0.41607, 3.14463, -0.784654, 2.56603, 0.569266, 14.4997, 18.2688, 2.13042, 4.72765, 30.598, 4.3042}

