

Supplementary file 4. nQuire histotest results for each strain analyzed in this work.

110_10

Diploid:

Norm SSR: 0.0552134

y-y slope: -0.13725, with std.Err: 0.0382047

r^2 : 0.179486

Triploid:

Norm SSR: 0.00637872

y-y slope: 0.442364, with std.Err: 0.0212552

r^2 : 0.880116

Tetraploid:

Norm SSR: 0.0206854

y-y slope: -0.556965, with std.Err: 0.0622311

r^2 : 0.575849

1282

Diploid:

Norm SSR: 0.0560525

y-y slope: -0.173305, with std.Err: 0.0200732

r^2 : 0.558186

Triploid:

Norm SSR: 0.0110048

y-y slope: 0.265766, with std.Err: 0.027109

r^2 : 0.619626

Tetraploid:

Norm SSR: 0.0106076

y-y slope: -0.042047, with std.Err: 0.0681987

r^2 : 0.00640143

14ANR23920

Diploid:

Norm SSR: 0.00796541

y-y slope: 0.575849, with std.Err: 0.0179947

r^2 : 0.945525

Triploid:

Norm SSR: 0.0462348

y-y slope: -0.352075, with std.Err: 0.102429

r^2 : 0.166841

Tetraploid:

Norm SSR: 0.0186045

y-y slope: 0.207154, with std.Err: 0.172576

r^2 : 0.0238392

9_16

Diploid:

Norm SSR: 0.00469496

y-y slope: 0.689673, with std.Err: 0.0188645

r^2 : 0.957724

Triploid:

Norm SSR: 0.0553508

y-y slope: -0.438739, with std.Err: 0.120707

r^2 : 0.182955

Tetraploid:

Norm SSR: 0.0240685

y-y slope: 0.231343, with std.Err: 0.205666

r^2 : 0.0209952

CBS180

Diploid:

Norm SSR: 0.00405022

y-y slope: 0.69373, with std.Err: 0.0111912

r^2 : 0.984878

Triploid:

Norm SSR: 0.0559558

y-y slope: -0.463219, with std.Err: 0.117936

r^2 : 0.207277

Tetraploid:

Norm SSR: 0.0240711

y-y slope: 0.210661, with std.Err: 0.204348

r^2 : 0.0176938

CH

Diploid:

Norm SSR: 0.00634326

y-y slope: 0.64761, with std.Err: 0.0241121

r^2 : 0.924395

Triploid:

Norm SSR: 0.051284

y-y slope: -0.376603, with std.Err: 0.117842

r^2 : 0.147562

Tetraploid:

Norm SSR: 0.0232117

y-y slope: 0.17607, with std.Err: 0.197344

r^2 : 0.0133122

CNM_CL6867

Diploid:

Norm SSR: 0.0522364

y-y slope: -0.131567, with std.Err: 0.020434

r²: 0.412677

Triploid:

Norm SSR: 0.0109143

y-y slope: 0.255593, with std.Err: 0.019971

r²: 0.735181

Tetraploid:

Norm SSR: 0.0111045

y-y slope: -0.104353, with std.Err: 0.0588597

r²: 0.0505799**IUM_96-0030**

Diploid:

Norm SSR: 0.0499344

y-y slope: -0.10266, with std.Err: 0.0231815

r²: 0.249477

Triploid:

Norm SSR: 0.00986203

y-y slope: 0.283758, with std.Err: 0.0123337

r²: 0.899713

Tetraploid:

Norm SSR: 0.013371

y-y slope: -0.248912, with std.Err: 0.0512343

r²: 0.285742**LL867**

Diploid:

Norm SSR: 0.056972

y-y slope: -0.17964, with std.Err: 0.0232311

r²: 0.503346

Triploid:

Norm SSR: 0.00827729

y-y slope: 0.348859, with std.Err: 0.0154679

r²: 0.896067

Tetraploid:

Norm SSR: 0.0145952

y-y slope: -0.271285, with std.Err: 0.0658031

r²: 0.223648

NRZ_BK_345

Diploid:

Norm SSR: 0.0507581

y-y slope: -0.117858, with std.Err: 0.0177256

r²: 0.428347

Triploid:

Norm SSR: 0.0109478

y-y slope: 0.244626, with std.Err: 0.0122517

r²: 0.871086

Tetraploid:

Norm SSR: 0.0124873

y-y slope: -0.217439, with std.Err: 0.0449416

r²: 0.284057

UCSC_1590

Diploid:

Norm SSR: 0.0584748

y-y slope: -0.189152, with std.Err: 0.0281812

r²: 0.43297

Triploid:

Norm SSR: 0.00782014

y-y slope: 0.380578, with std.Err: 0.022632

r²: 0.827372

Tetraploid:

Norm SSR: 0.0168089

y-y slope: -0.365686, with std.Err: 0.0701593

r²: 0.315285