Plot 38: mean fitness (t−test)
Group 237: retromer complex (N=6) Benschop

Y-axis: bir1−17 37 SDM_rhl_CNGHT MDRMDP DIL bir1−17 MS 2010

X-axis: cSGA 37 SDM_rhl_CNGT MDRMDP DIL cSGA MS 2009

Labels: MUK1, VPS35, PEP8, VPS29, VPS17, VPS5