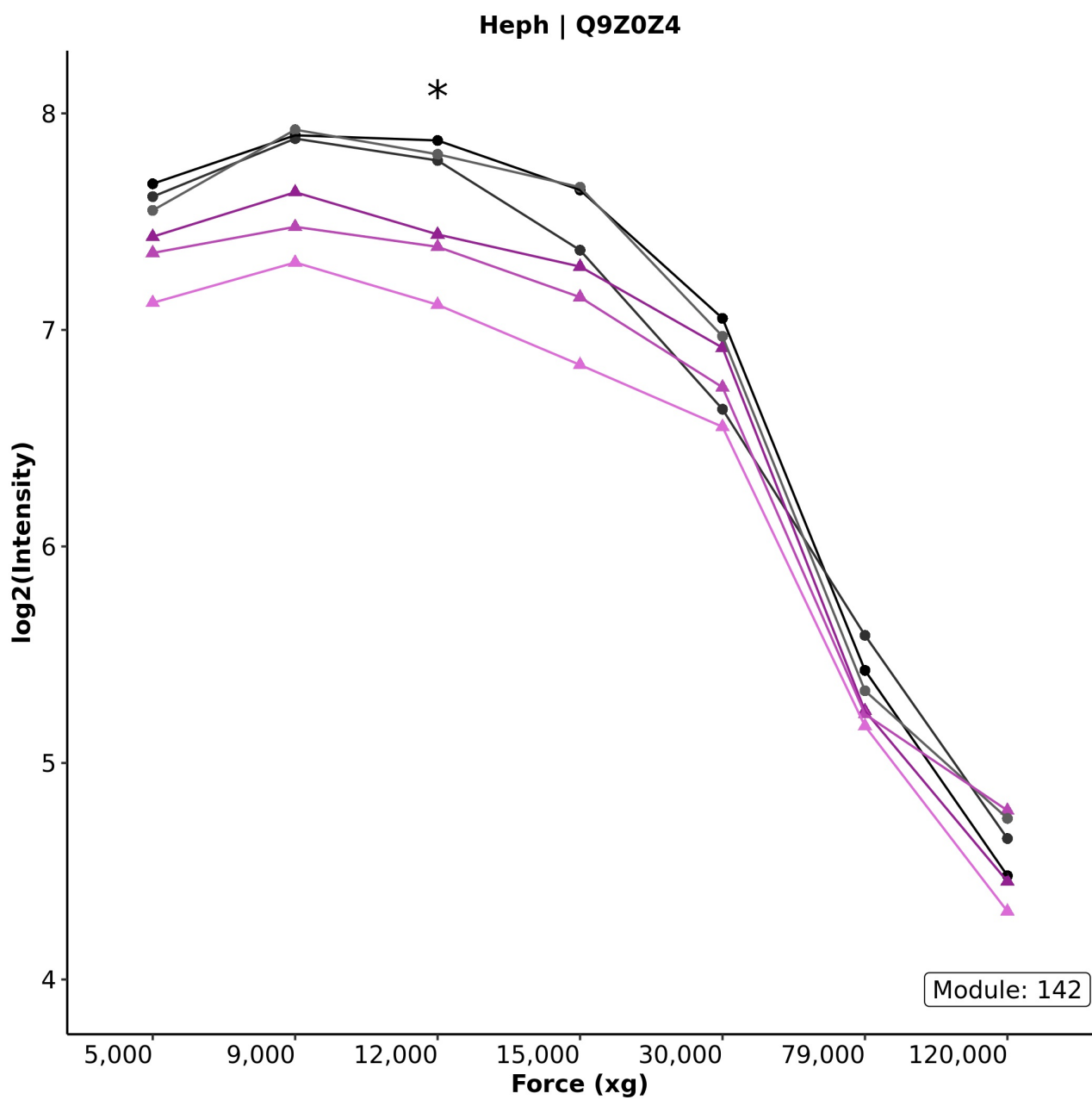


Supplementary File 4

Plots of protein abundance measured in seven subcellular fractions for **85** proteins with an intrafraction significant difference between WT and MUT groups (FDR < 0.1). The 7 subcellular fractions coorespond to fractions 4-10 in Geledaki *et al.*, 2019 [37].

Example Plot:



Plot Attributes:

Attribute	Description
x-axis	Centrifugal force (xg) used to obtain the cooresponding subcellular fraction.
y-axis	Log2(Normalized Protein Intensity)
point shape	Circles and triangles indicate animal genotype, WT and MUT, respectively.

line color Lines connecting points are colored according to biological replicate.

Statistical Significance:

Symbol Threshold

.	FDR < 0.1
*	FDR < 0.05
**	FDR < 0.005
***	FDR < 0.0005