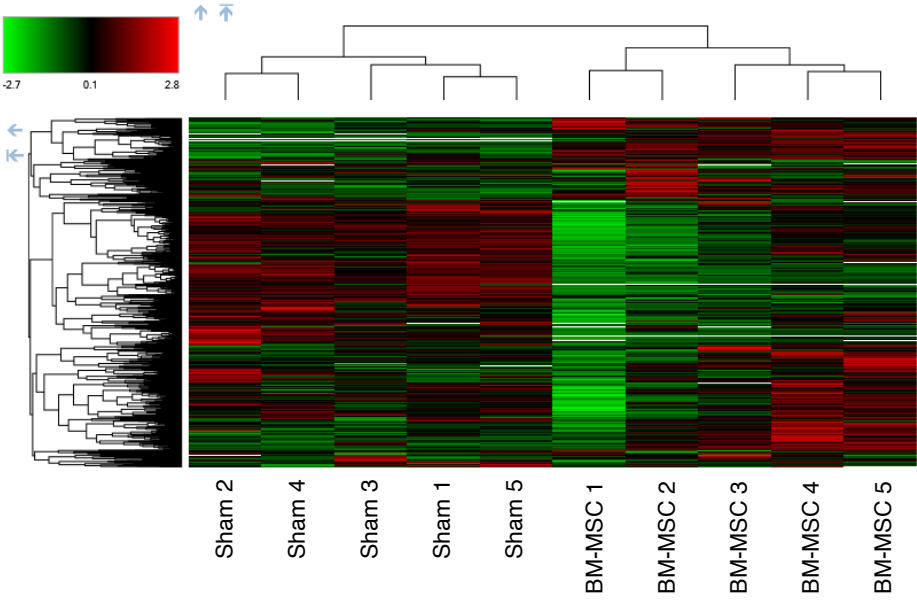
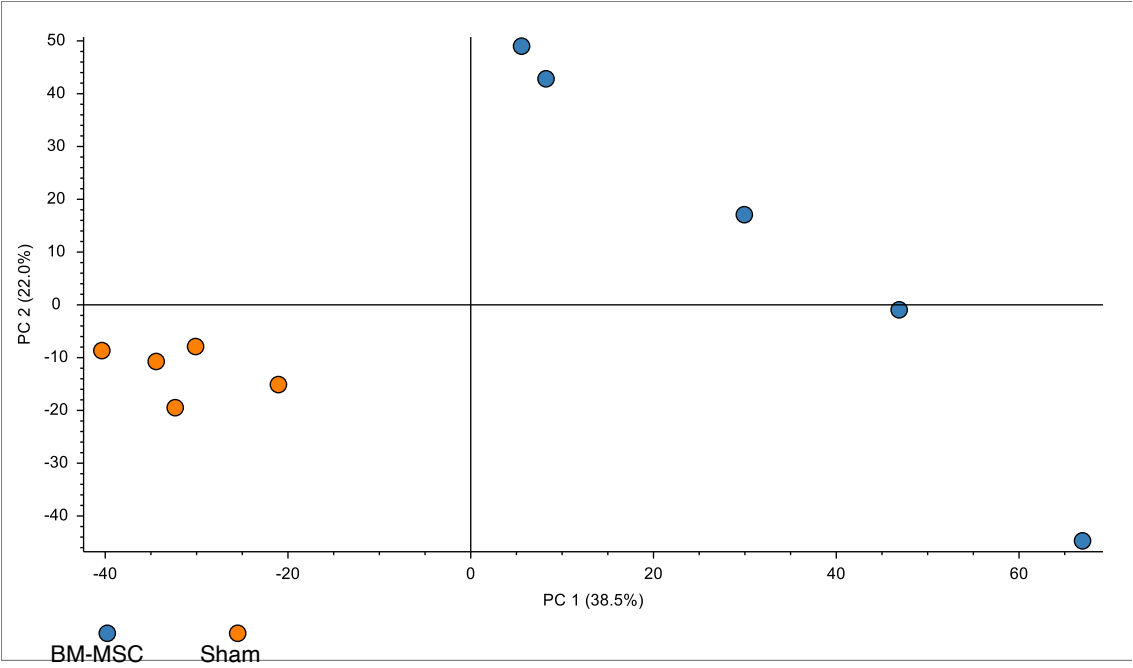


**FIGURE S2**

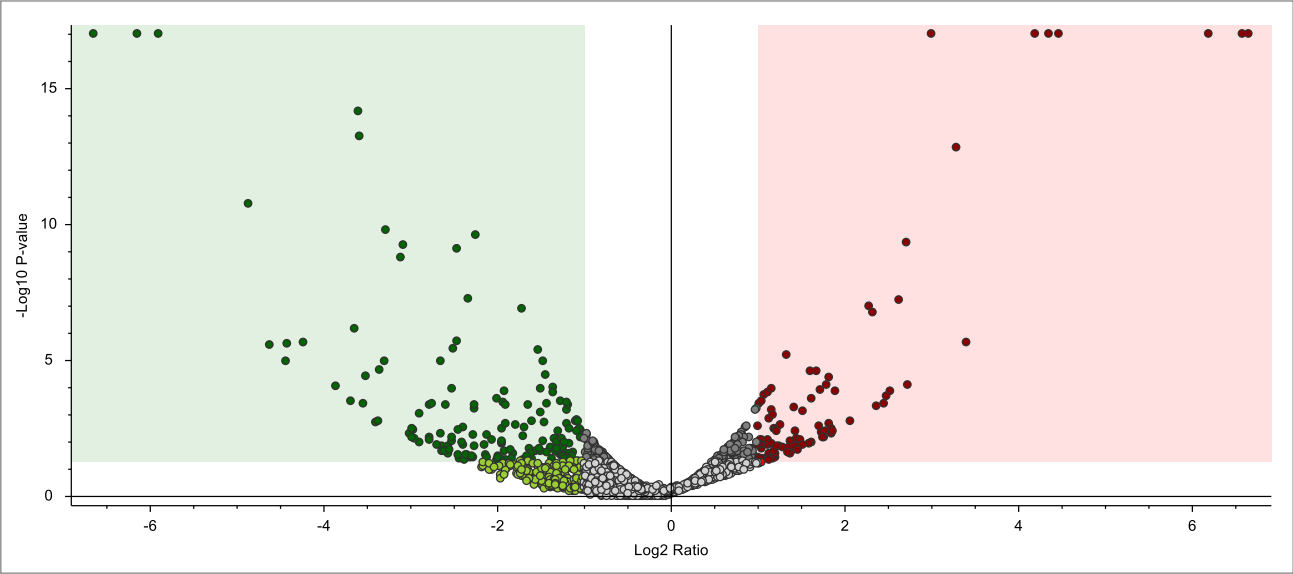
Data Source: Proteins : Abundances (Normalized)  
Distance Function: Pearson  
Linkage Method: Complete  
Scaling: Scale Before Clustering



**a**



**b**



**c**