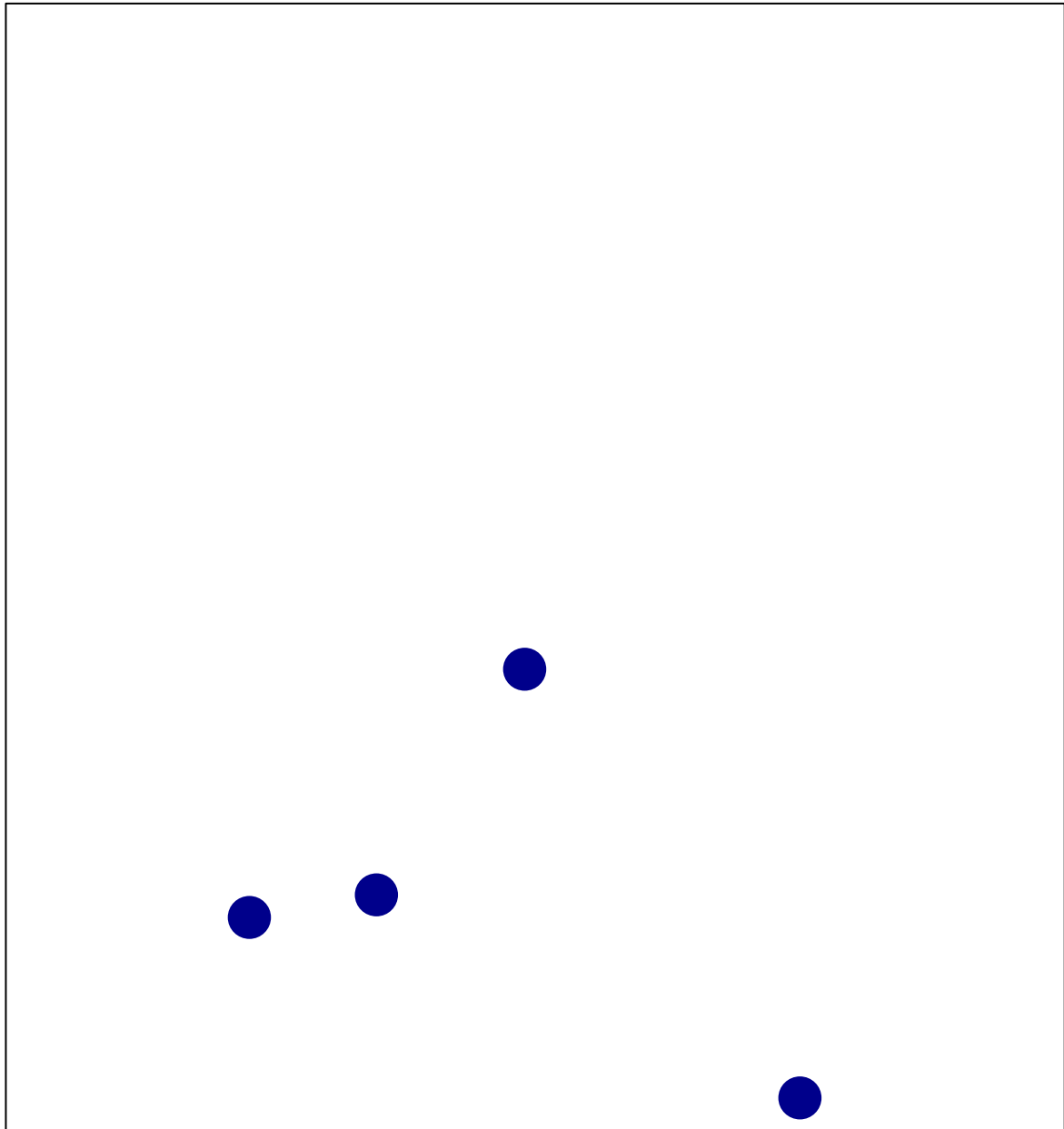


# WEBER\_METHYLATED\_ICP\_IN\_SPERM\_UP



# features = 4  
chi-square p = 0.85

# WEBER\_METHYLATED\_ICP\_IN\_SPERM\_UP



# features = 4 , max = 1