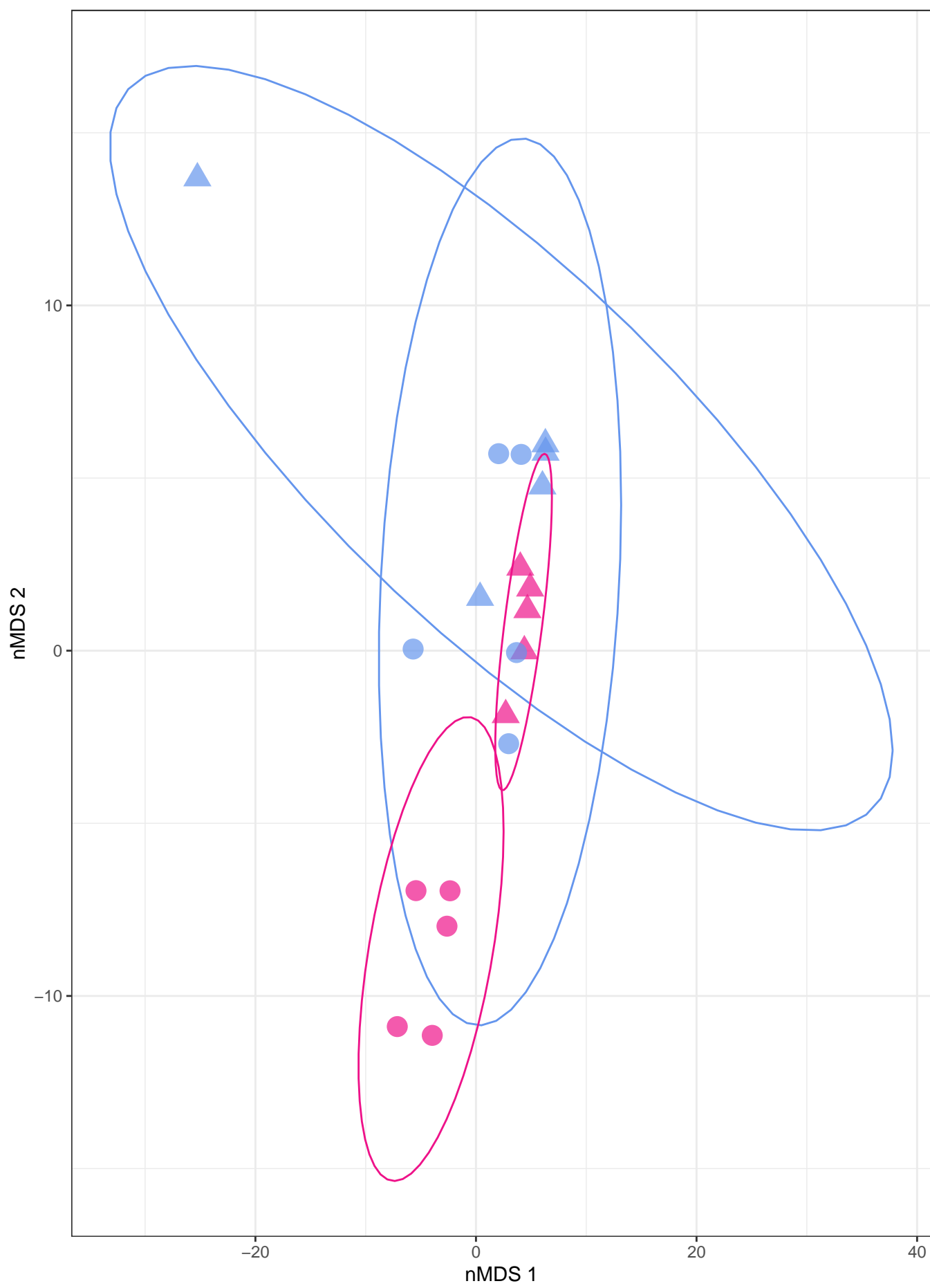A) All gene regions

B) Significant DMG