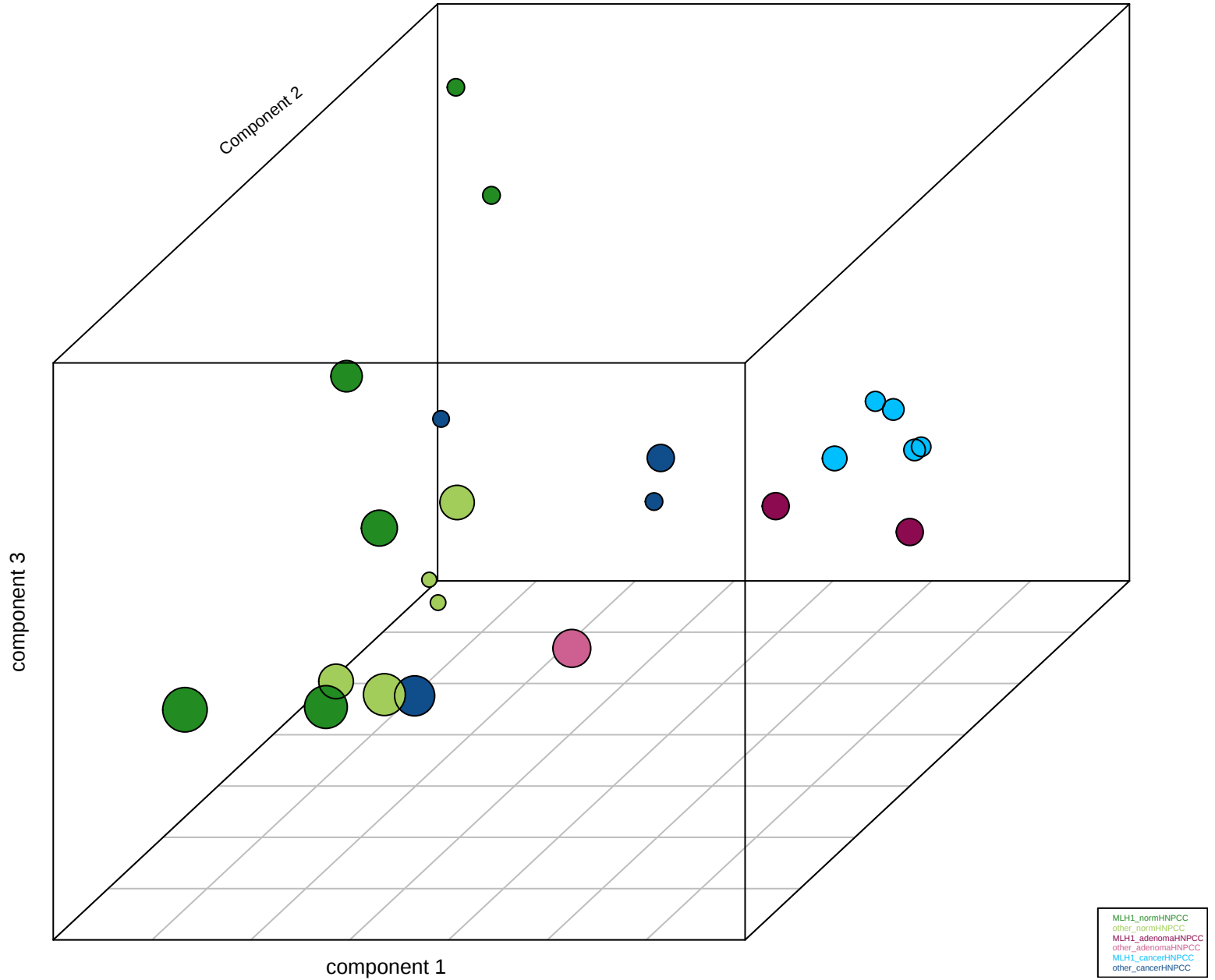
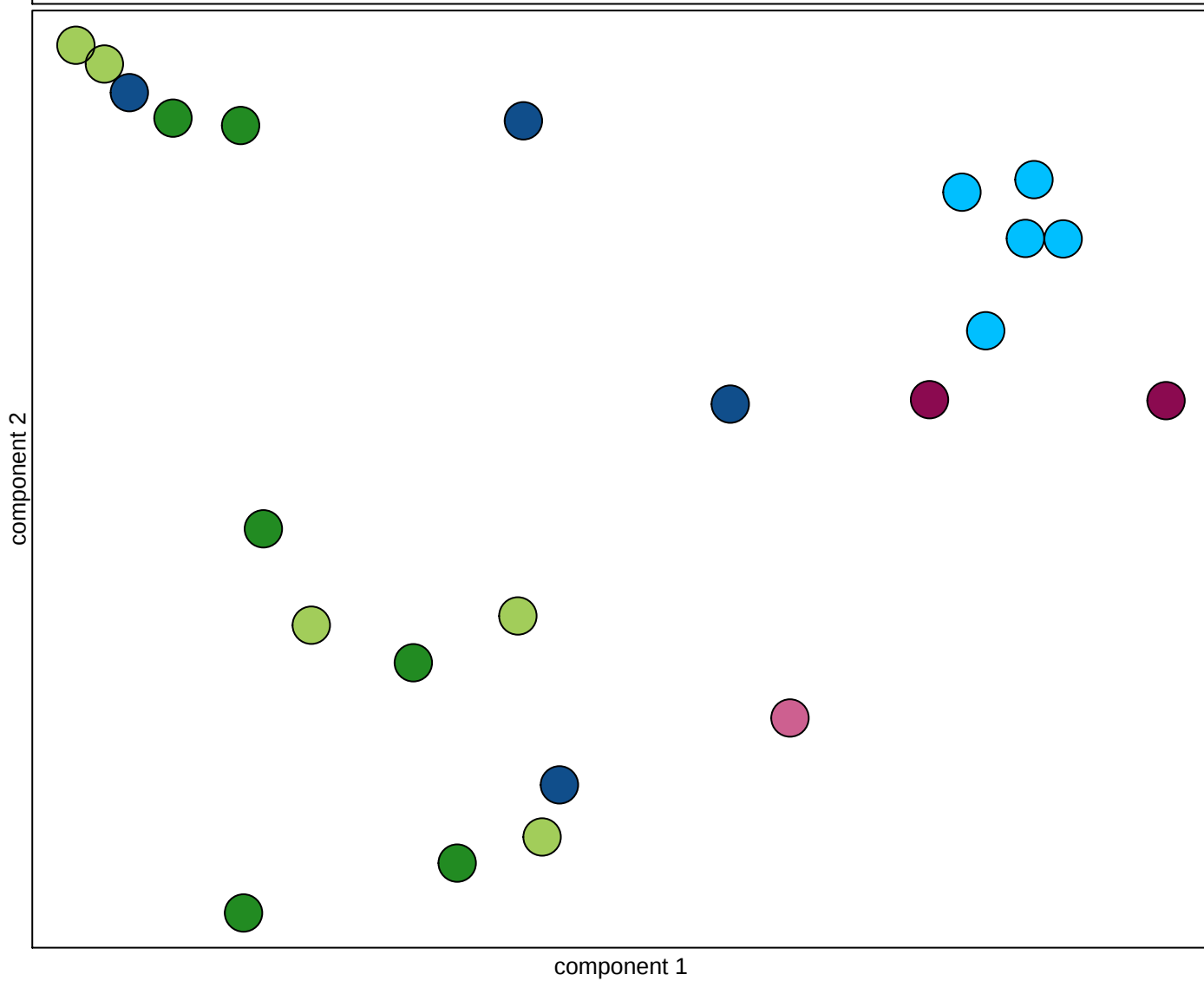
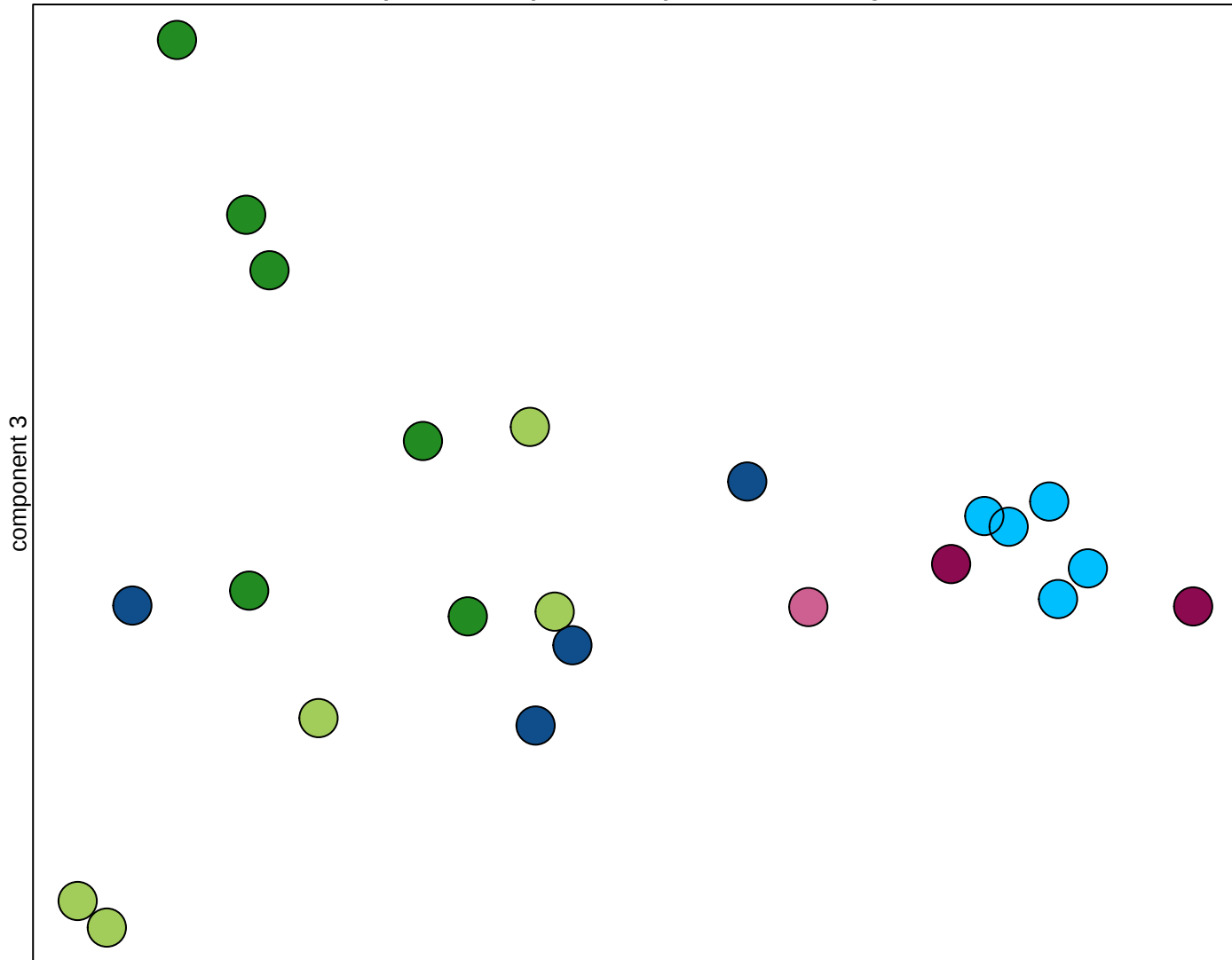


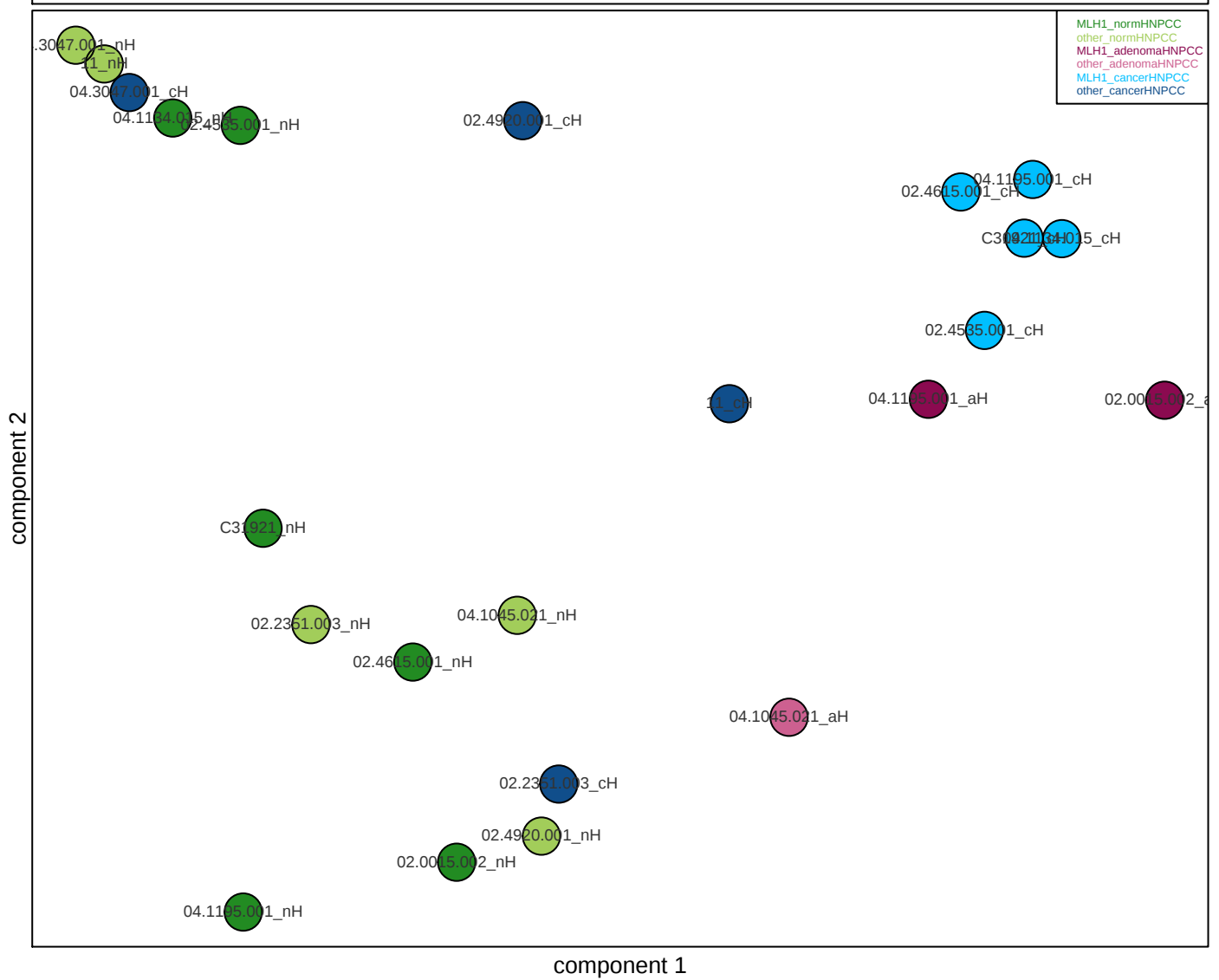
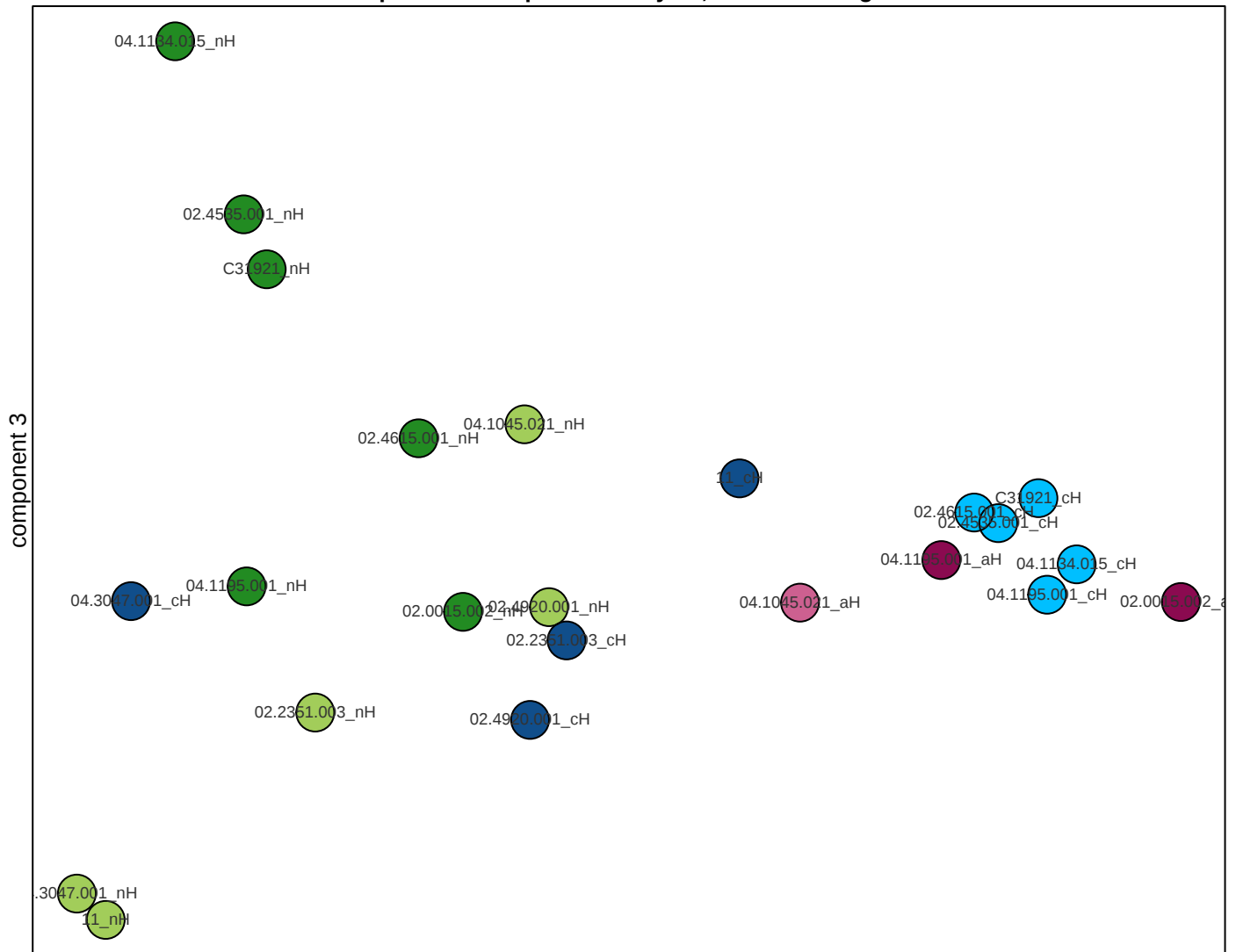
Independent component analysis, all 1600 Metagenes



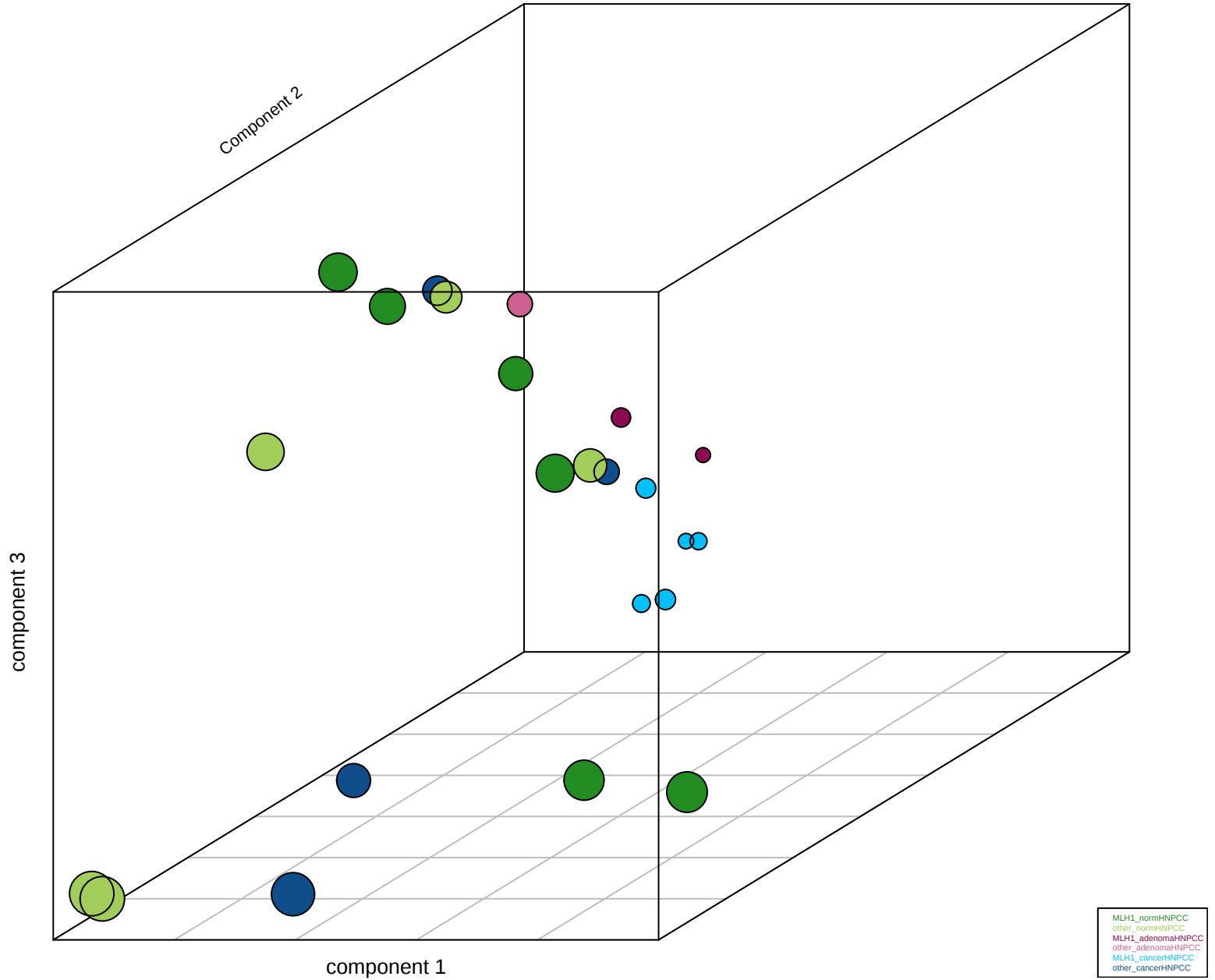
Independent component analysis, all 1600 Metagenes



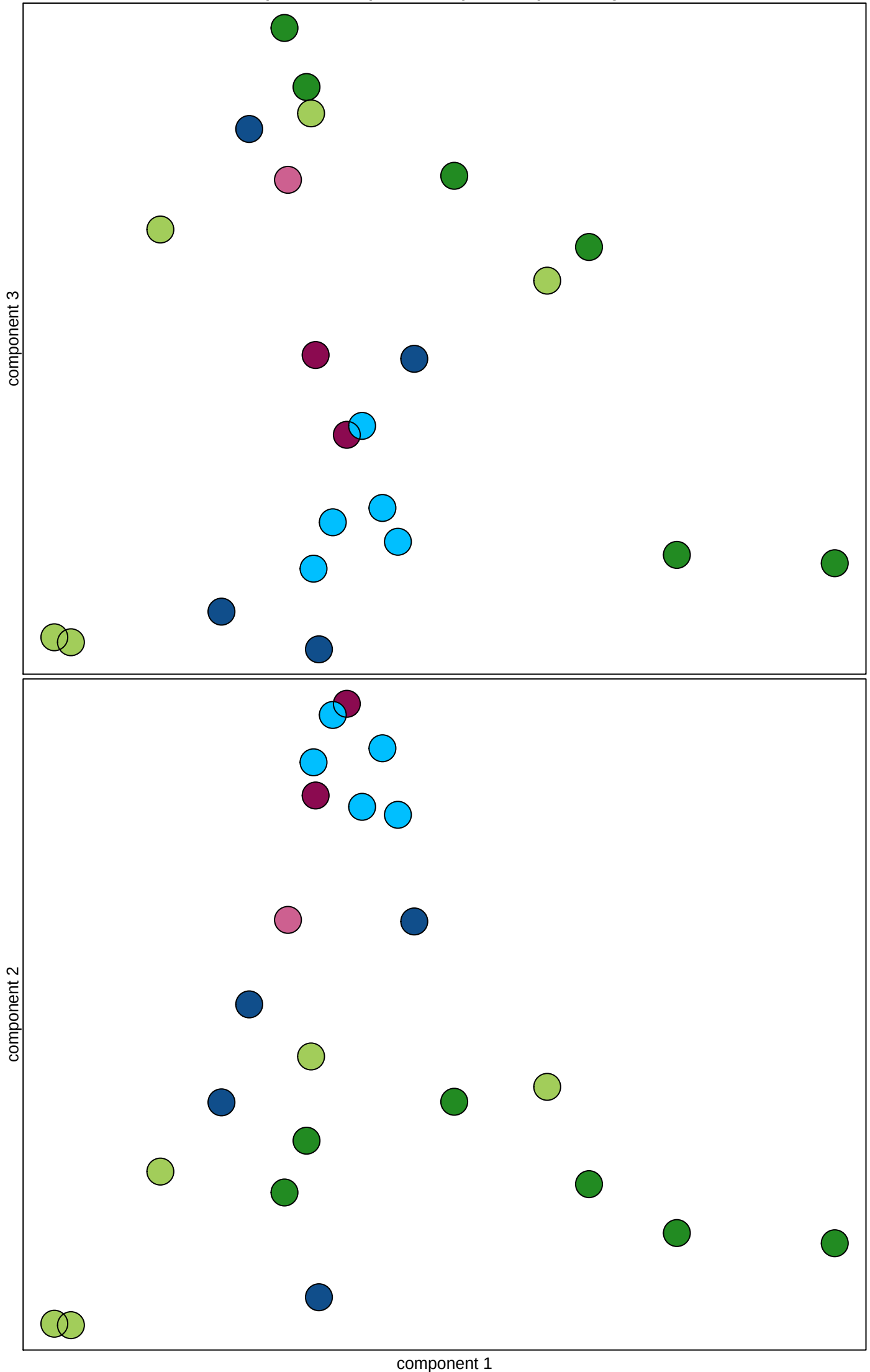
Independent component analysis, all 1600 Metagenes



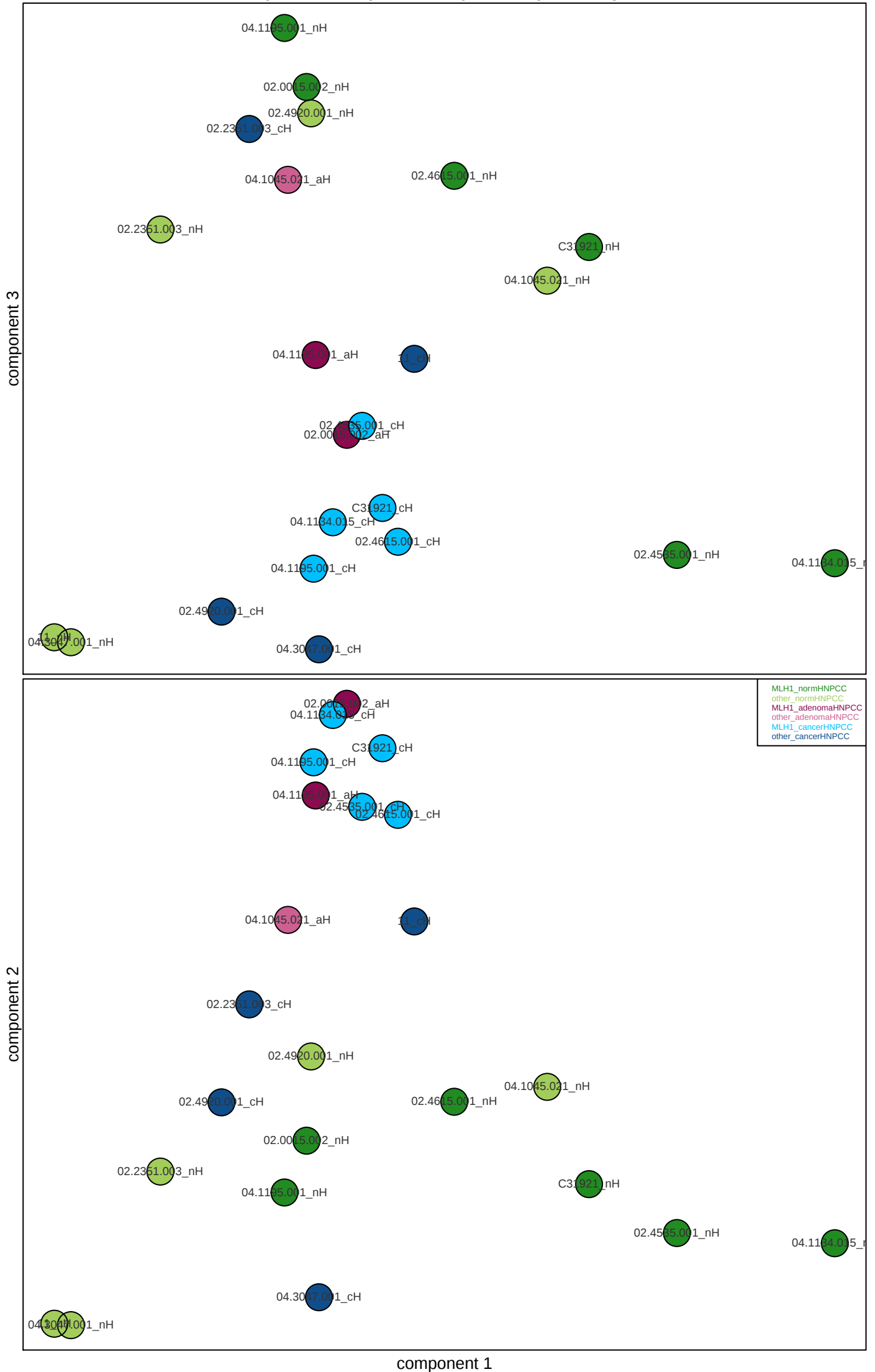
Independent component analysis, 33 Spot-Metagenes



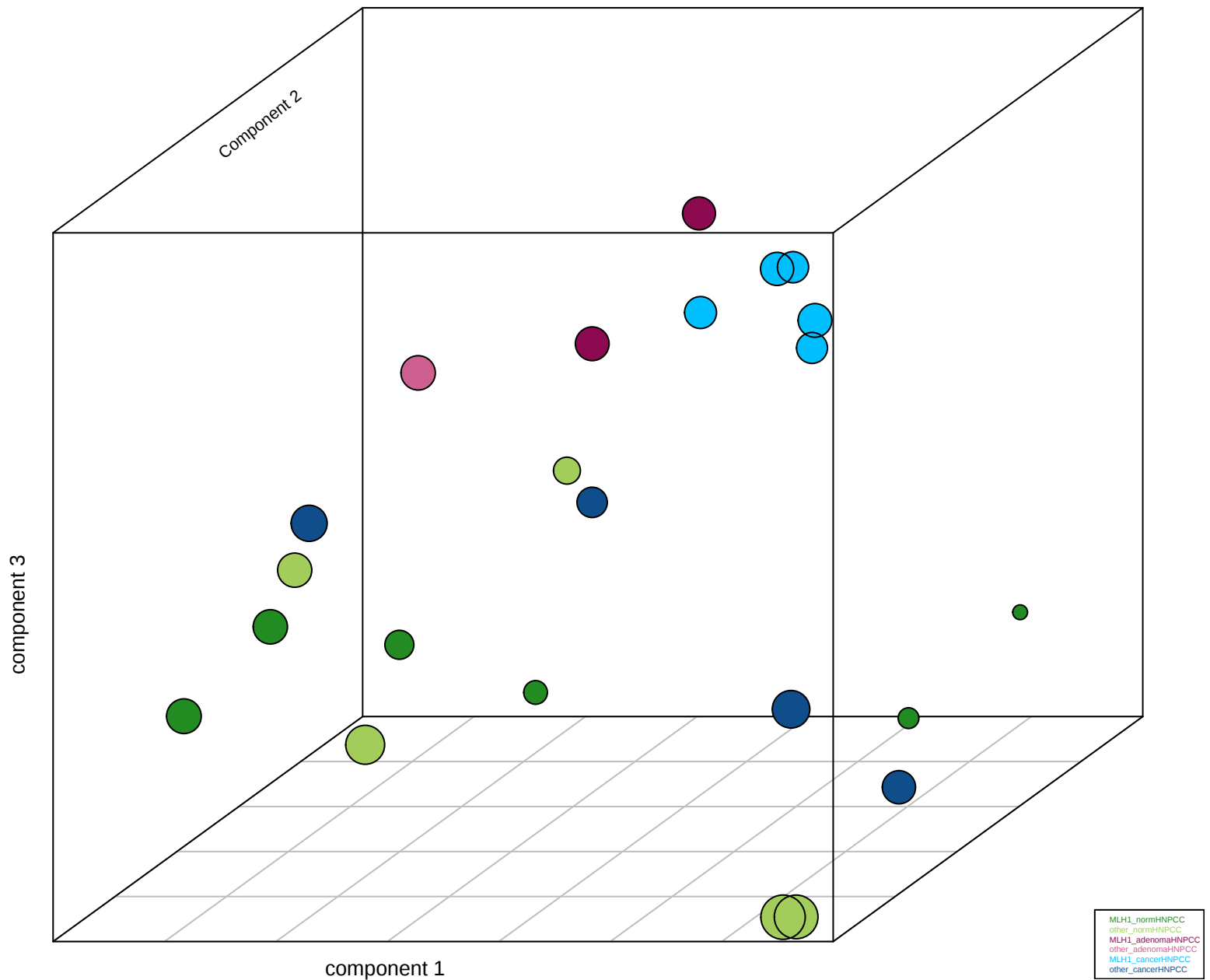
Independent component analysis, 33 Spot-Metagenes



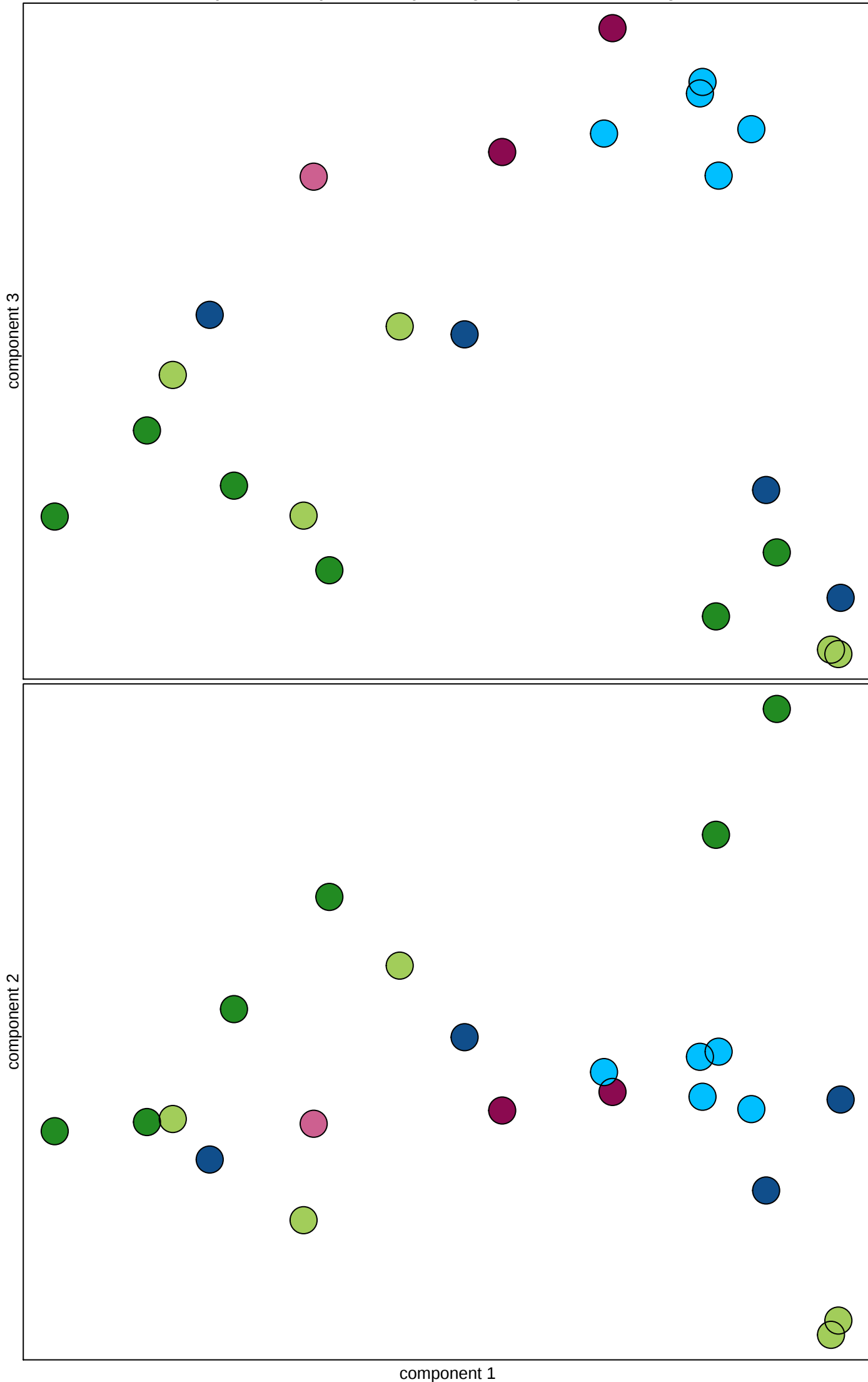
Independent component analysis, 33 Spot-Metagenes



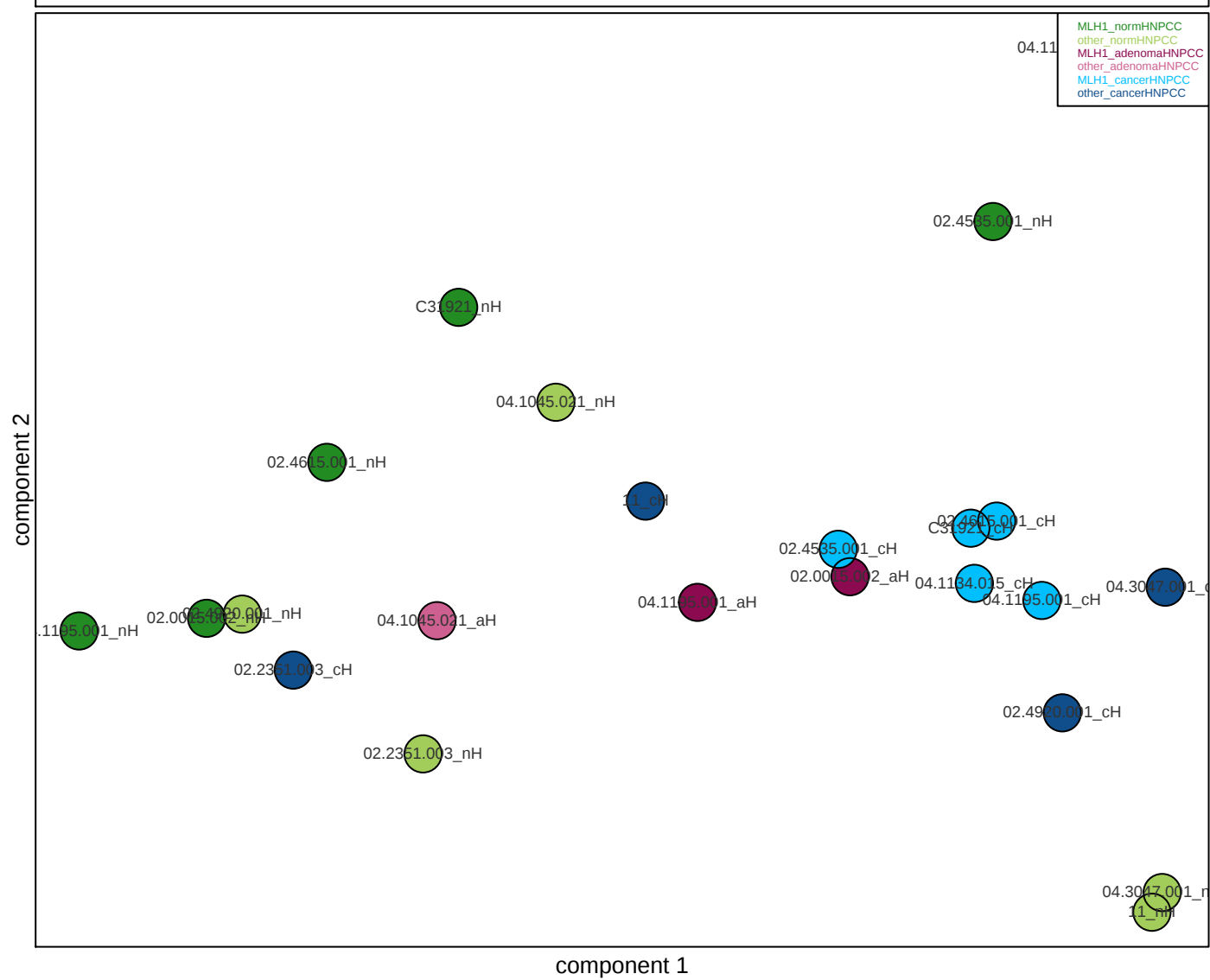
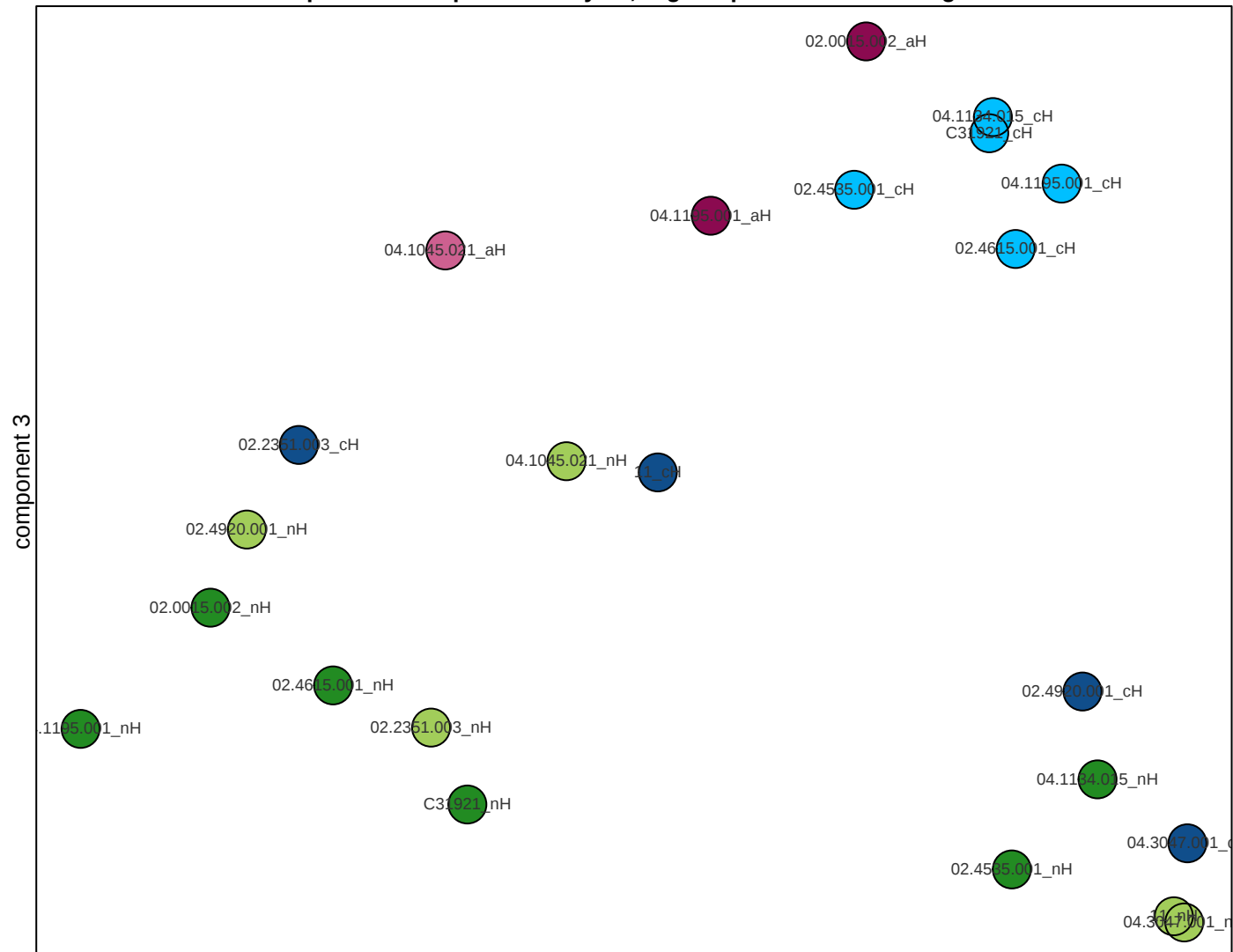
Independent component analysis, High Expression: 100 Metagenes



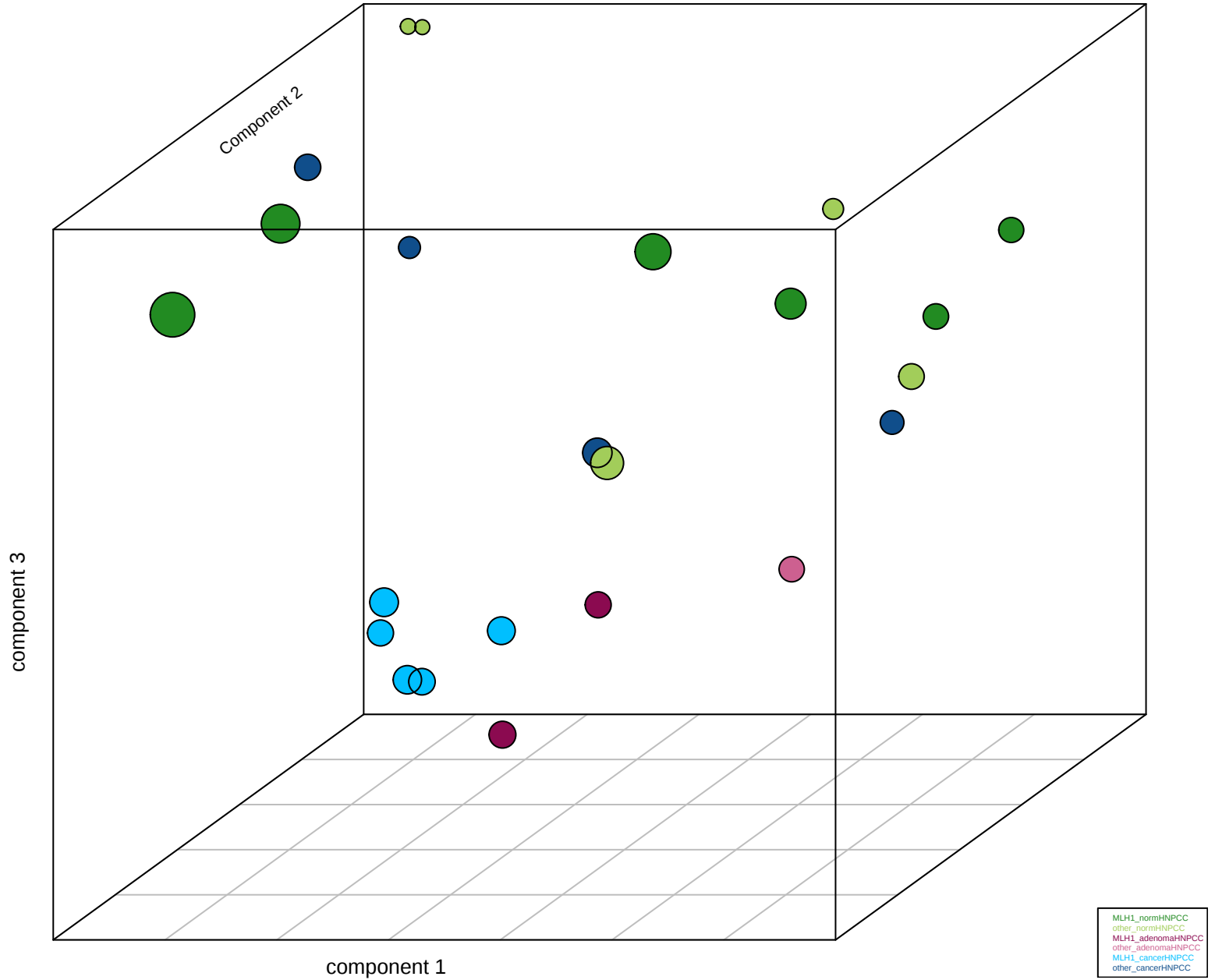
Independent component analysis, High Expression: 100 Metagenes



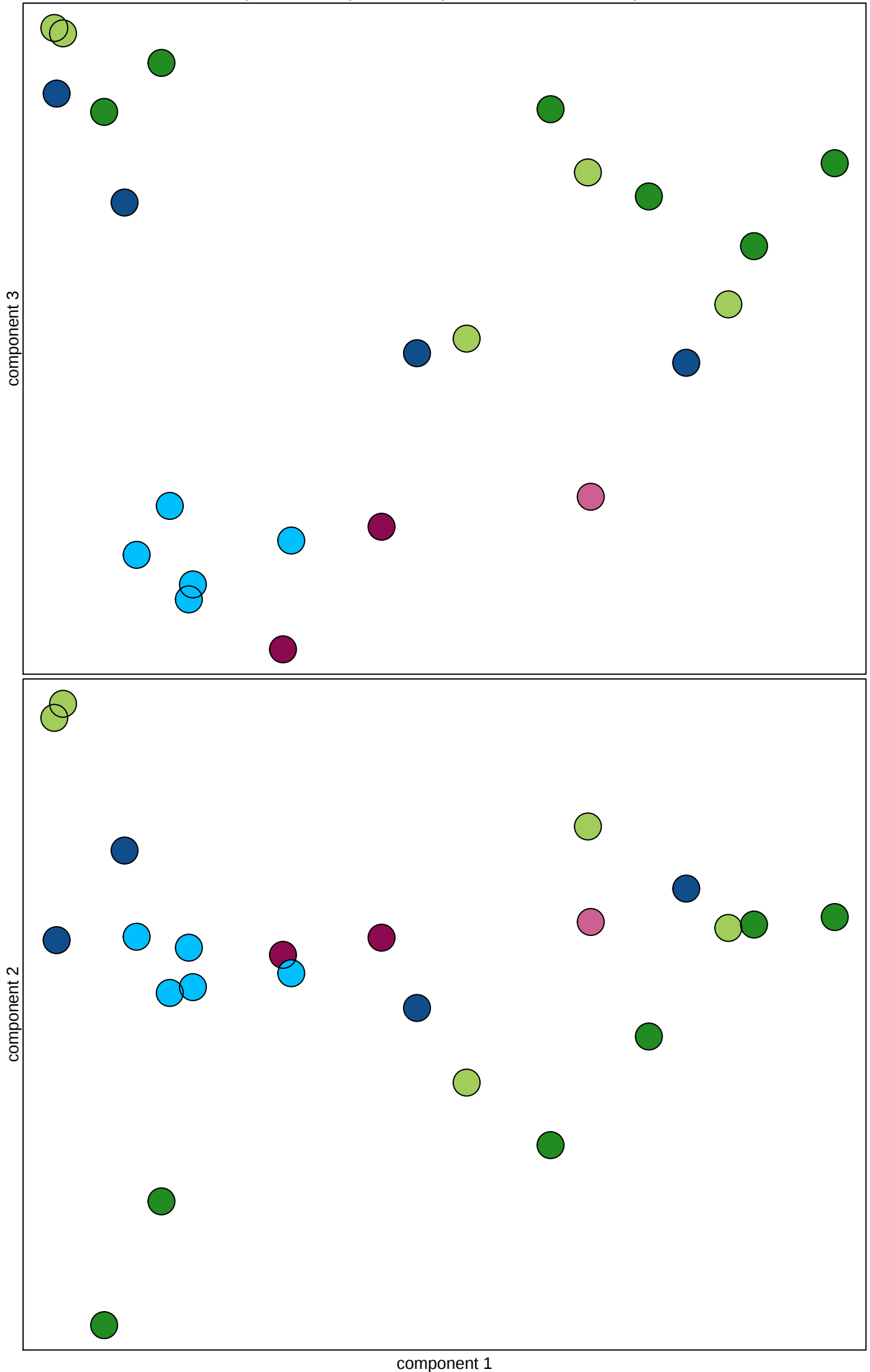
Independent component analysis, High Expression: 100 Metagenes



Independent component analysis, Variance: 100 Metagenes



Independent component analysis, Variance: 100 Metagenes



Independent component analysis, Variance: 100 Metagenes

