

bisulfite sequencing method originally developed by AI CINA sark anages in Australia [32, 33].

Aim 2: To determine patterns of methylation in different physiological contexts.

A) We already know that individual ^{13}C isotope ratios reflectively indicate of selected MTag.

commercially (on F13Choice[®] catalogue). We can measure liver, skeletal muscle, heart

the work properly. For instance, efforts to establish a culture model systems will not need to be