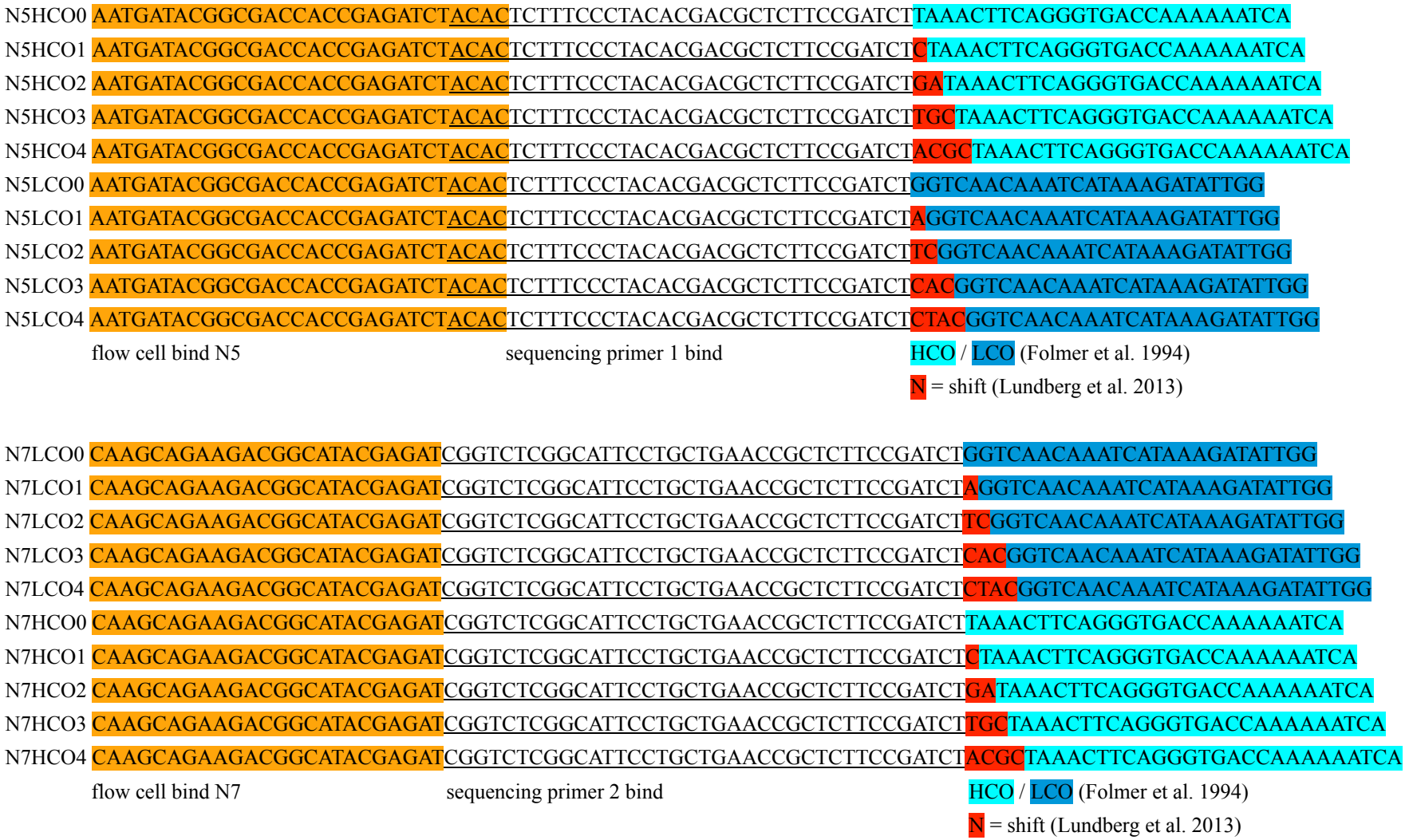

Figure S1. Fusion COI Primers developed in this study. The primers are based on the Folmer et al. (1994) primers. They include MiSeq flow cell and sequencing primer binding regions. All primers amplify an 832-bp fragment that can be sequenced directly after purification. Up to 10 libraries can be uniquely tagged and pooled in one MiSeq run with the tags consisting of the HCO or LCO primer + the primer shift up to 4 bp (always 4 bp in total). It is recommended that all 10 primer pairs are used in the following combination to maximise sequence diversity:

N5HCO0+N7LCO4, N5HCO1+N7LCO3, N5HCO2+N7LCO2, N5HCO3+N7LCO1, N5HCO4+N7LCO0,
N5LCO0+N7HCO4, N5LCO1+N7HCO3, N5LCO2+N7HCO2, N5LCO3+N7HCO1, N5LCO4+N7HCO0