

## The sequences of staggered primers

### Primers for profiling the V3-V4 region of 16S DNA

#### Primers for reaction 1 (direct orientation of the amplicon)

TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGCCTACGGGAGGCAGCAG  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGgaCCTACGGGAGGCAGCAG  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGtagCCTACGGGAGGCAGCAG  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGagcaattCCTACGGGAGGCAGCAG  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGGGACTACHVGGGTWTCTAAT  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGcaGGACTACHVGGGTWTCTAAT  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGtctGGACTACHVGGGTWTCTAAT  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGatctGGACTACHVGGGTWTCTAAT

#### Primers for reaction 2 (reverse orientation of the amplicon)

TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGGGACTACHVGGGTWTCTAAT  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGcaGGACTACHVGGGTWTCTAAT  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGatctGGACTACHVGGGTWTCTAAT  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGtctactGGACTACHVGGGTWTCTAAT  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGCCTACGGGAGGCAGCAG  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGgaCCTACGGGAGGCAGCAG  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGtagCCTACGGGAGGCAGCAG  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGagcaattCCTACGGGAGGCAGCAG

### Primers for subtyping *Blastocystis* sp.

#### Reaction 1 (direct orientation of the amplicon)

TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGGGAGGTAGTGACAATAAATC  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGaGGAGGTAGTGACAATAAATC  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGcaGGAGGTAGTGACAATAAATC  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGactGGAGGTAGTGACAATAAATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGTGCTTTTCGCACTTGTTTCATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGaTGCTTTTCGCACTTGTTTCATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGcaTGCTTTTCGCACTTGTTTCATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGactTGCTTTTCGCACTTGTTTCATC

#### Reaction 2 (reverse orientation of the amplicon)

TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGTGCTTTTCGCACTTGTTTCATC  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGaTGCTTTTCGCACTTGTTTCATC  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGcaTGCTTTTCGCACTTGTTTCATC  
TCGTCGGCAGCGTCAGATGTGTATAAGAGACAGactTGCTTTTCGCACTTGTTTCATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGGGAGGTAGTGACAATAAATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGaGGAGGTAGTGACAATAAATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGcaGGAGGTAGTGACAATAAATC  
GTCTCGTGGGCTCGGAGATGTGTATAAGAGACAGactGGAGGTAGTGACAATAAATC

## *Legend:*

### Illumina overhang tags:

plain uppercase

TCGTCGGCAGCGTC (P5-tag, a Read1 Nextera pre-adapter) and GTCTCGTGGGCTCGG (P7-tag, a Read 2 Nextera pre-adapter). The P5- and P7-tags are the sites where second-round indexing primers anneal.

### Illumina pad sequence:

underlined

AGATGTGTATAAGAGACAG; it is the annealing site of the Illumina sequencing primers. The read 1 and read 2 thus both start from the specific locus primer (or the heterogeneity spacer if it is present). The Illumina overhang tag along with the pad sequence make together the whole overhang sequence - these are the annealing sites for the read 1 sequencing primer (the P5-tag and the following pad) and the read 2 sequencing primer (the P7-tag and the sequencing primer).

### Heterogeneity spacers (primer staggering, heterogeneity spacers):

ga, agt... (lowercase, yellow)

### Specific locus primers:

bold uppercase

16S rDNA profiling: **CCTACGGGAGGCAGCAG, GGACTACHVGGGTWTCTAAT** (bold uppercase): specific portions of primers V3-forward (341-F, [1]) and V4-reverse (806-R, [2]).

*Blastocystis* subtyping: **GGAGGTAGTGACAATAAATC, TGCTTTCGCACTTGTTTCATC** are specific portions of F (ILMN\_Blast505\_532F) and R (ILMN\_Blast998\_1017R) primers for subtyping of *Blastocystis* sp. according to Maloney et al [3]

## **References:**

- [1] Muyzer G, de Waal EC, Uitterlinden AG (1993) Profiling of complex microbial populations by denaturing gradient gel electrophoresis analysis of polymerase chain reaction-amplified genes coding for 16S rRNA. Appl Environ Microbiol 59: 695-700
- [2] Caporaso JG, Lauber CL, Walters WA, et al. (2011) Global patterns of 16S rRNA diversity at a depth of millions of sequences per sample. Proc Natl Acad Sci U S A 108 Suppl 1: 4516-4522
- [3] Maloney JG, Molokin A, Santin M (2019) Next generation amplicon sequencing improves detection of *Blastocystis* mixed subtype infections. Infect Genet Evol 73: 119-125