

Online Resource 3

Article title: Genomic prediction of starch content and chipping quality in tetraploid potato using genotyping-by-sequencing

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Authors: Elsa Sverrisdóttir, Stephen Byrne, Ea Høegh Riis Sundmark, Heidi Øllegaard Johnsen, Hanne Grethe Kirk, Torben Asp, Luc Janss, and Kåre L. Nielsen

Corresponding author: Elsa Sverrisdóttir, Aalborg University, Department of Chemistry and Bioscience, Fredrik Bajers Vej 7H, 9220 Aalborg, Email: esv@bio.aau.dk; Telephone number: +45 5055 3092

Adapter and primer sequences

5' Index adapter

- 1) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTTGCA
- 2) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTTGCGA
- 3) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTCGCTT
- 4) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTGGTTGT
- 5) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTCCAGCT
- 6) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTTATTTT
- 7) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTTTGCTT
- 8) 5'-ACACTCTTTCCCTACACGACGCTCTTCCGATCTACGACTAC

5' Index adapter reverse complement

- 1) 5'-CWGTGCAAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT
- 2) 5'-CWGTCGCAAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT
- 3) 5'-CWGAAGCGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT
- 4) 5'-CWGACAACAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT
- 5) 5'-CWGAGCTGGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT
- 6) 5'-CWGAAAAATAAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT
- 7) 5'-CWGAAGCAAGAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT
- 8) 5'-CWGGTAGTCGTAGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT

3' Index adapter

- A) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACATCACGATCTCGTATGCCGTCTTCTGCTTG
- B) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACCGATGTATCTCGTATGCCGTCTTCTGCTTG
- C) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACTTAGGCATCTCGTATGCCGTCTTCTGCTTG
- D) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACTGACCAATCTCGTATGCCGTCTTCTGCTTG
- E) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACACAGTGATCTCGTATGCCGTCTTCTGCTTG
- F) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACGCCAATATCTCGTATGCCGTCTTCTGCTTG
- G) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACAGTCAAATCTCGTATGCCGTCTTCTGCTTG
- H) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACACTTGAATCTCGTATGCCGTCTTCTGCTTG
- I) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACGATCAGATCTCGTATGCCGTCTTCTGCTTG
- J) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACTAGCTTATCTCGTATGCCGTCTTCTGCTTG
- K) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACGGCTACATCTCGTATGCCGTCTTCTGCTTG
- L) 5'-CWG AGATCGGAAGAGCACACGTCTGAACTCCAGTCACCTTGTAATCTCGTATGCCGTCTTCTGCTTG

3' Index adapter reverse complement

- A) 5'-CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- B) 5'-CAAGCAGAAGACGGCATACGAGATACATCGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- C) 5'-CAAGCAGAAGACGGCATACGAGATGCTTAAGTACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- D) 5'-CAAGCAGAAGACGGCATACGAGATTGGTCAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- E) 5'-CAAGCAGAAGACGGCATACGAGATCACTGTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- F) 5'-CAAGCAGAAGACGGCATACGAGATATTGGCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT

- G) 5'-CAAGCAGAAGACGGCATAACGAGATTTGACTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- H) 5'-CAAGCAGAAGACGGCATAACGAGATCAAGTGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- I) 5'-CAAGCAGAAGACGGCATAACGAGATCTGATCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- J) 5'-CAAGCAGAAGACGGCATAACGAGATAAGCTAGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- K) 5'-CAAGCAGAAGACGGCATAACGAGATGTAGCCGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT
- L) 5'-CAAGCAGAAGACGGCATAACGAGATTACAAGGTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT

PCR primer 1

5'-AATGATACGGCGACCAACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT

PCR primer 2

5'-CAAGCAGAAGACGGCATAACGAGAT