Table S1. The primers designed for gene cloning, expression, deletion, complementation and mutant identification.

Primers	Sequences (5′-3′) *	Purpose
Sec22-F	ATGCTCTGCGCCTCAGTCGACGACG	Cloning *BbSec22*
Sec22-R	TTAGAAAATTCGCCAGTAGATGAAG	
Sec22q-F	CGAGATGAGCAGCAGGTTGAG	qRT-PCR for R-SNAREs
Sec22q-R	AATTCGCCAGTAGATGAAGAGGAG	
Ykt6 q-F	GCGAAACAACTATGGCGAATT	
Ykt6 q-F	ACCTTGCTGAGGACCTGATGG	
Nyv1q-F	CGTGCCCTTTGGCTACCTTGT	
Nyv1q-R	AGTGTCTTGAGCGTGCCGTTG	
Nyv2q-F	CACCCAAATCACCCATCTGCC	
Nyv2q-R	GCTATACTCGACCATGAGCGTCTT	
Snc1q-F	CAGGCTCTGCAAGCTCAAATCG	
Snc1q-R	CGCATCTTCATGTCCTTCCACC	
18S-F	TGGTTTCTAGGACCGCCGTAA	
18S-R	CCTTGGCAAATGCTTTCGC	
Sec22up-F	CCCAAGCTTCGCTGTTGAAATGTGCTCTGTCGGT	Deleting *BbSec22*
Sec22up-R	CGGCGAGCTCATGACGGCATGTTTCAATGATGGGA	
Sec22dn-F	GGAAGATCTGAACGGCGTACGGAAACGAATGCAC	
Sec22dn-R	CTAGACTAGTGAATGAACGGGAAACTCATGGATGG	
Sec22c-F	GGGGACAAGTTTGTACAAAAAAGCAGGCTGTGGT GTAACCTTGCGATGCTCAG	Rescuing *BbSec22*
Sec22c-R	GGGGACCACTTTGTACAAGAAAGCTGGGTCCCTG GACCACTTTCTTGTTTCTTC	
Id-F	AGAACCAGGTCCGCCAAATC	PCR, RT-PCR for *BbSec22*
Id-R	GAGGAGAATAATGAAGCCCAGCAC	

* The introduced cleavage sites of the enzymes *Hin*dIII, *Sac*I, *Bgl*II and *Spe*I are underlined.