

GSBRNA2T00134741001	ATGATCAAACGAAACCGGGGAGAACCCCTGCTGTGATGGCCGGGAAGACGATGTGACGGAATCATCCCTTCTCAGAAACAAAAGCTGGCGAAGAAGTGTGT	98
Bo1015899	ATGATCAAACGAAACCGGGGAGAACCCCTGCTGTGATGGCCGGGAAGACGATGTGACGGAATCATCCCTTCTCAGAAACAAAAGCTGGCGAAGAAGTGTGT	100
Consensus	atgatcaaacgaaacccgggagaaaccccgcggtgatggccgggaagacgatgtgacggaatcatcccttctcagaaacaaaagccggcggaagaagtgtgt	
GSBRNA2T00134741001	GAGAACCACCCTTCTGGAGTCAAAGGCCAAACITTCGGAAGACTTAACCGGAATAATCCGTGTGATGGGAACAAGTCGGCGAAAACCTCCCGGAAAGA	198
Bo1015899	GAGAACCACCCTTCTGGAGTCAAAGGCCGAACITTCGGAAGACTTAACCGGAATAATCCGTGTGATGGGAACAAGTCGGCGAAAACCTCCCTGAAAGA	200
Consensus	gagaaccacccttctggagtcaaaagccaaaccttcgaaagacttaaccggaataatccggtgatgggaacaagtcggcgaaaaactccccgaaaga	
GSBRNA2T00134741001	TTGAATTTTGATAAGAAGAAGAGAAATTGAGAAAGAGAGAAGCCGACACAATCATCAAACCAAGTTTCCTTTTCGTTATGATAAAAAGTTATTTACTTTA	298
Bo1015899	TTGAATTTTGATAAGAAGAAGAGAAATTGAGAAAGAGAGAAGCCGACACAATCATCAAACCAAGTTTCCTTTTCGTTATGATAAAAAGTTATTTACTTTA	300
Consensus	ttgaattttgataagaagaagagaaattgagaaagagagaagccgacacaatcatcaaaaccaagtttcttttcgttatgataaaaagttatttacttta	
GSBRNA2T00134741001	TTTATAATTTACTATATTTAAAACCTTTAAAAGTGTATGTTTTACATTCACACGACCGAAACAGAGAGCAGCCGACTCTTTCATTTGGCGCGCTGAGCTA	398
Bo1015899	TTTATAATTTACTATATTTAAAACCTTTAAAAGTGTATGTTTTACATTCACACGACCGAAACAGAGAGCAGCCGACTCTTTCATTTGGCGCGCTGAGCTA	400
Consensus	tttataatttactatattttaaacttttaaagtgatgttttacattcacacgaccgaaacagagagcagccgactctttcatttggcggcgctgagcta	
GSBRNA2T00134741001	AGAAAAGGGTTTTACACGATAAAAAATATTTCTATATCTATCAGCTTAAACCTTATTTCTTTGTCTTAAGAACATCTGATAAACTCTGTCTCTCTCAACAG	498
Bo1015899	AGAAAAGGGTTTTACACGATAAAAAATATTTCTATATCTATCAGCTTAAACCTTATTTCTTTGTCTTAAGAACATCTGATAAACTCTGTCTCTCTCAACAG	500
Consensus	agaaaagggttttacacgataaaaaatatttctatatctatcagcttaaaccttatctttgtcttaagaacatctgataaaactctgtctctctcaacag	
GSBRNA2T00134741001	AGATTTCTTTCCCTCTAAGAAAAATAATAATTTCTTTTTCTTTTTTATTTGTGGGTTTATAAGCTTCAAGTTTCAGACAAGGATTATAG.....	589
Bo1015899	AGATTTCTTTCCCTCTAAGAAAAATAATAATTTCTTTTTCTTTTTTATTTGTGGGTTTATAAGCTTCAAGTTTCAGACAAGGATTATAGGGAGAGAAT	600
Consensus	agatctctttccctctaagaaaaataataatcttttcttttttatttgtgggtttataagcttcaagtttcagacaaggattatagggagagaat	
GSBRNA2T00134741001	..	589
Bo1015899	TG	602
Consensus	tg	