

Sequence Listing

1	Sequence Listing Information	
1-1	File Name	sequence listing.xml
1-2	DTD Version	V1_3
1-3	Software Name	WIPO Sequence
1-4	Software Version	2.3.0
1-5	Production Date	2024-01-11
2	General Information	
2-1	Current application: IP Office	
2-2	Current application: Application number	
2-3	Current application: Filing date	
2-4	Current application: Applicant file reference	0073
2-5	Earliest priority application: IP Office	
2-6	Earliest priority application: Application number	
2-7	Earliest priority application: Filing date	
2-8zh	Applicant name	JIANGXI FUSHINE BIOTECHNOLOGY CO., LTD
2-8	Applicant name: Name Latin	Jiangxi Fuxiang Biotechnology Co., Ltd
2-9	Inventor name	
2-9	Inventor name: Name Latin	
2-10zh	Invention title	USE OF FUSARIUM BRACHYGIBBOSUM IN

		PRODUCTION OF MYCELIAL PROTEIN
2-11	Sequence Total Quantity	6
3-1	Sequences	
3-1-1	Sequence Number [ID]	1
3-1-2	Molecule Type	DNA
3-1-3	Length	20
3-1-4-1	Features Location/Qualifiers	source 1..20 mol_type= other DNA organism= synthetic construct
3-1-5	Residues	atgggtaagg argacaagac 20
3-2	Sequences	
3-2-1	Sequence Number [ID]	2
3-2-2	Molecule Type	DNA
3-2-3	Length	19
3-2-4-1	Features Location/Qualifiers	source 1..19 mol_type= other DNA organism= synthetic construct
3-2-5	Residues	ggargtacca gtsatcatg 19
3-3	Sequences	
3-3-1	Sequence Number [ID]	3
3-3-2	Molecule Type	DNA
3-3-3	Length	22

3-3-4-1	Features Location/Qualifiers	source 1..22 mol_type= other DNA organism= synthetic construct
3-3-5	Residues	ggaagtaaaa gtcgtaacaa gg 22
3-4	Sequences	
3-4-1	Sequence Number [ID]	4
3-4-2	Molecule Type	DNA
3-4-3	Length	19
3-4-4-1	Features Location/Qualifiers	source 1..19 mol_type= other DNA organism= synthetic construct
3-4-5	Residues	ggtccgtgtt tcaagacgg 19
3-5	Sequences	
3-5-1	Sequence Number [ID]	5
3-5-2	Molecule Type	DNA
3-5-3	Length	612
3-5-4-1	Features Location/Qualifiers	source 1..612 mol_type= other DNA organism= synthetic construct
3-5-5	Residues	gactctggca agtcgaccac tgtaagtacc accaagcctc gacccgcct agatttgggtg 60 gggtagtctc aagattcatc catactgaca tactttgata gaccggtcac ttgatctacc 120 agtgcggtgg tatcgacaag cgaaccateg

		agaagttcga gaaggttggt ctcatttccc 180 tegatcgcgcc gcccttttac ccatcgatcc atcattegac tcgtctctc acgacgactc 240 gatecgcgcc cgttaccccg ctcgagctca aaaattttgc gaatcaatcg taattttttt 300 cttgatgggg ctcatacccc gccactegag tgacgggcgc gcttgccctc tteccacagt 360 catattcata tgggcgcgca tcatcacgtg tctaccagtc actaaccact tgacaatagg 420 aagccgccga gctcggtaag ggttccttca agtacgectg ggttcttgac aagetcaagg 480 ccgagcgtga gcgtggtatc accattgata tcgtctctg gaagttcgag actcctcgtc 540 actatgtcac cgtcattggg atgctgtcac cgttgccctc atcacatcct catactaaca 600 cgtccaccag ac 612
3-6	Sequences	
3-6-1	Sequence Number [ID]	6
3-6-2	Molecule Type	DNA
3-6-3	Length	1065
3-6-4-1	Features Location/Qualifiers	source 1..1065 mol_type= other DNA organism= synthetic construct
3-6-5	Residues	ccgagtttac aactcccaaa cccctgtgaa cataccttta tgttgccctg gcggatcagc 60 ccgcgccccg taaaacggga cggccccgcg caggaaccac aaaactctga ttttagtgta 120 acttctgagt ctaaaaaaca aataaatcaa aactttcaac aacggatetc ttggttctgg 180 catcgatgaa gaacgcagca aaatgcgata

agtaatgtga attgcagaat tcagtgaate 240
 atcgaatctt tgaacgcaca ttgcgcccgc
 cagtattctg gcgggcatgc ctgttcgagc 300
 gtcatttcaa cectcaagcc cccgggttg
 gtgttgggga tcgggctgta ctccagcccg 360
 gccccgaaat ctagtggcgg tctcgtgca
 gcctccattg cgtagtagct aacacctgc 420
 aactggaacg cggcgcggcc aagccgtaa
 acccccaact tctgaatgtt gacctcgga 480
 caggtaggaa taccgctga acttaagcat
 atcaataagc ggaggaaaag aaaccaacag 540
 ggattgccct agtaacggcg agtgaagcgg
 caacagctca aatttgaaat ctggctctcg 600
 ggcccagatt gtaatttgta gaggatgctt
 ttgatgcggt gccttcgag ttcctggaa 660
 cgggacgcca tagagggtga gagccccgc
 tggttgatg ccaaattctt gtaaagctcc 720
 ttcgacgagt cgagtagttt gggaatgctg
 ctctaaatgg gaggtatatg tcttctaaag 780
 ctaaataccg gccagagacc gatagcgcac
 aagtagagtg atcgaaagat gaaaagcact 840
 ttgaaaagag agttaaaaag tacgtgaaat
 tgttgaaagg gaagcgttta tgaccagact 900
 tgggcttggg taatcatctg gggttctccc
 cagtgcactt ttccagtcca ggccagcatc 960
 agttttcgcc gggggataaa ggttcggga
 atgtggctcc ctccggggag tgttatagcc 1020
 cgttgcgtaa taccctggcg gggactgagg
 ttcgcgttc tgcaa 1065