[CLAIMS]

[Claim 1]
A polynucleotide having promoter activity, wherein at least one nucleotide selected from the group consisting of the 92nd, 94th, 102nd, 103rd, 249th, and 251st nucleotides of the polynucleotide sequence of SEQ ID NO: 1 or 2 is substituted with a different nucleotide,

wherein the substitution of the 92nd nucleotide is substitution to cytosine(C);
the substitution of the 94th nucleotide is substitution to cytosine(C);
the substitution of the 102nd nucleotide is substitution to cytosine(C);
the substitution of the 103rd nucleotide is substitution to thymine(T);
the substitution of the 249th nucleotide is substitution to adenine(A); and
the substitution of the 251st nucleotide is substitution to thymine(T).

[Claim 2]
The polynucleotide of claim 1, wherein the 249th and 251st nucleotides of the polynucleotide having promoter activity are substituted with different nucleotides.

[Claim 3]
The polynucleotide of claim 1, wherein the 92nd, 94th, 102nd, and 103rd nucleotides of the polynucleotide having promoter activity are substituted with different nucleotides.

[Claim 4]
The polynucleotide of claim 1, wherein the 92nd, 94th, 102nd, 103rd, 249th, and 251st nucleotides of the polynucleotide having promoter activity are substituted with different nucleotides.

[Claim 5]
The polynucleotide of claim 1, wherein the polynucleotide having promoter activity consists of one polynucleotide sequence selected from SEQ ID NOS: 3 to 32.

[Claim 6]
A composition for expressing a gene comprising the polynucleotide of any one of claims 1 to 5.

[Claim 7]
A vector comprising the polynucleotide of any one of claims 1 to 5 and a gene encoding a target protein.

[Claim 8]
The vector of claim 7, wherein the target protein has glutamate–cysteine ligase activity.

[Claim 9]
A microorganism belonging to the genus *Saccharomyces* sp. comprising the polynucleotide of any one of claims 1 to 5, a polynucleotide comprising the polynucleotide and a gene encoding a target protein, or a vector including the same.

[Claim 10]
The microorganism of claim 9, wherein the target protein is a polypeptide having glutamate–cysteine ligase activity.

[Claim 11]
A method of producing glutathione comprising culturing the microorganism of claim 9 in a culture medium.

[Claim 12]
The method of claim 11, further comprising recovering glutathione from at least one selected from the cultured microorganism, a dried product of the microorganism, an extract of the microorganism, a culture product of the microorganism, and a lysate of the microorganism.

[Claim 13]
A use of a polynucleotide sequence, wherein at least one nucleotide selected from the group consisting of the 92nd, 94th, 102nd, 103rd, 249th, and 251st nucleotides of the polynucleotide sequence of SEQ ID NO: 1 or 2 is substituted with a different nucleotide, as a promoter, wherein

the substitution of the 92nd nucleotide is substitution to cytosine(C);
the substitution of the 94th nucleotide is substitution to cytosine(C);
the substitution of the 102nd nucleotide is substitution to cytosine(C);
the substitution of the 103rd nucleotide is substitution to thymine(T);
the substitution of the 249th nucleotide is substitution to adenine(A); and
the substitution of the 251st nucleotide is substitution to thymine(T).