



Investigation of taste-altering mechanism: A model Study for sweet taste providing activity of miracle fruit

味覚改変作用機構の解明:ミラクリンフルーツの甘味提供機構をモデルとして

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Abstract

Miraculin (MIR) is a kind of taste-modifying glycoprotein, which can modify a sour taste into a sweet taste, isolated from a red fruit called Richadella dulcifica, also named as miracle fruits, a native shrub in tropical West Africa. It is a homodimer that consists of 2 glycosylated 191-amino acid polypeptides that are cross-linked by disulfide bonds. MIR has unique taste-modifying properties. Though flat in taste at neutral pH, it shows taste-modifying activity to convert sourness to sweetness at acidic pH. Although this interesting sensory effect has been previously characterized, the molecular mechanism underlying the taste-modifying action of MIR is unknown.





<u>Introduction</u>

The position of S - S bonds:
1C47-C92, 2C152-C155,
3C148-C159, 4C138-C138(subunit)
The basic character of the MIR monomer:

191 amino acid residues,

Molecular weight is 24600Da (13.9% of sugars)

The dimer can modify a sour taste to a sweet taste, but the monomer can not.

S.Theerasilp, Y. Kurihara (1988 J. Biol. Chem. 263: 11536–9)

ATGANGGANT TANCANIGCT CICCICCIC TICTICG CONTROL STORM KELL TWELL S SERVED TO SALL A A A SERVED TO SALL A A SERVED TO SALL A SERVED

The amino acid sequence of miraculin

Trp100

Cys152

Cys152

Cys155

Cys159

Cys138

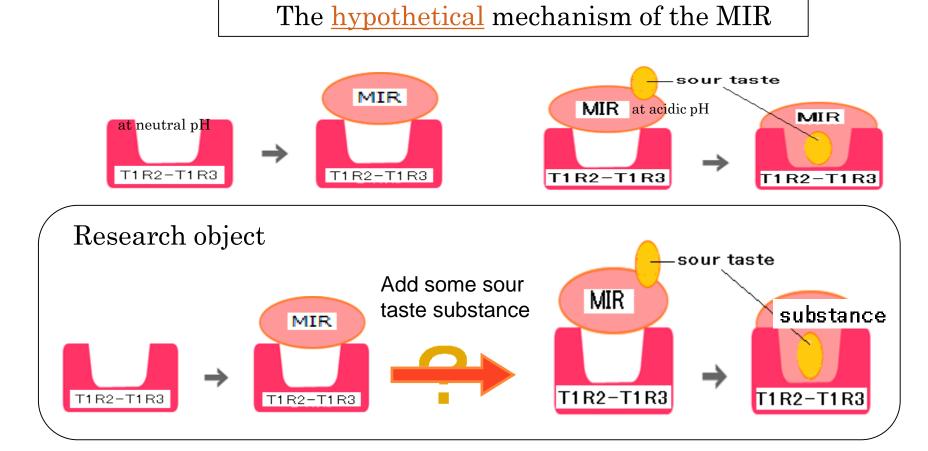
Trp131

f MIR monomor 3D structure, by Discovery Studio

Photo of miracle fruits

Simulation model of MIR monomer 3D structure by Discovery Studio® 2.5 (Template model: Kunitz (STI) type inhibitor, 1R8N)

mechanism of taste primary taste sweet taste, bitter taste, sour taste, salty taste, umami taste umami taste gustatory nerve Type II GABA GABA GABA GABA GABA Type III GABA GABA GABA GABA GABA GABA GABA Type III GABA GABA Type III GABA Type III GABA GABA Type III Type III GABA Type III GABA Type III Type



Method and result

1. Sensory evaluation: We got the results by method, with sensory evaluation the crude extract from the miracle fruits.

Confirmation of the activity of miraculin

	Sour taste	Sweet taste
Ascorbic acid	-	++
Citric acid	+	+++
Acetic acid	-	+
Malic acid	-	+
Lactic acid	_	_

The PSE (Point of Subjective Equality) of the 4 kinds of the organic acid composition, when we can feel sweet taste.

	クエン酸		リンゴ酸		酢	酸	アスコルビン酸		
甘味を呈する範囲	下限	上限	下限	上限	下限	上限	下限	上限	
濃度 % (g/v)	0.01	0.05	0.008	0.038	0.007	0.037	0.021	0.103	
mol濃度 (mmol/L)	0.52	2.60 0.60		2.83	1.17	6.16	1.19	5.85	
рН	3.42	2.96	3.48	3.07	3.87 3.48		3.63	3.23	
pKa	3.15 , 4.77 , 6.40		3.4, 5.1		4.7	76	4.17		

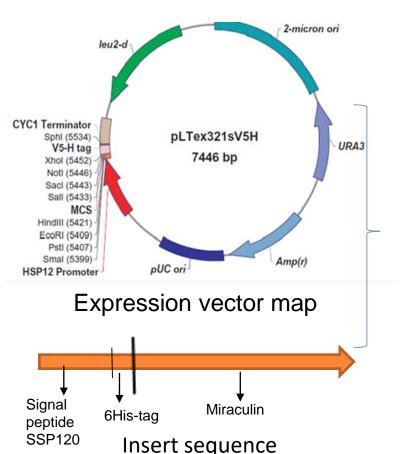
which sample can make you feel strongest sweet taste?

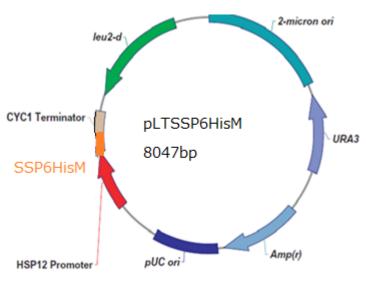
which sample can make you feel stronger sweet taste between 2 different concentration compared with 0.8% sucrose?

パネル	試料									
	0.01%クエン酸	0.008%リンゴ酸	0.007%酢酸							
А	1	3	2							
В	2	1	3							
C	1	2	3							
D	3	1	2							
E	1	3	2							
F	3	2	1							
合計	11	12	13							

	試料				試料		試料					試料			
パネル	0.01% クエン 酸	0.05% クエン 酸	0.8%ス クロー ス	パネル	0.008% リンゴ 酸	0.038% リンゴ 酸	0.8%ス クロー ス	パネル	0.007% 酢酸	0.037% 酢酸	0.8%スクロース	パネル	0.021 %アス コルビ	0.103% アスコ ルビン	0.8% スク ロー
Α	2	3	1	Α	2	3	1	A	1	3	2		ン酸	酸	ス
В	3	1	2	В	3	1	2	В	3	1	2	A	3	2	1
C	2	3	1				2	С	3	2	1	В	3	1	2
D	3	1	2	С	2	3	1	D	1	2	3	С	2	1	3
	3	1	2	D	3	1	2	Г	1	2	2	D	2	1	3
	3	1	_	Е	1	3	2	t	1	3	2	Е	2	3	1
F	2	1	3	F	2	3	1	F	1	3	2	F	3	1	2
G	2	3	1		_	3	1	G	2	3	1	G	2	3	1
Н	3	2	1	G	2	1	3	Н	2	2	1	Н	3	2	1
1	2	3	1	Н	2	3	1	П	3	2	1	- 1	3	2	1
J	2	1	3	1	2	1	3	I	1	3	2	J	1	3	2
K	3	1	2	J	2	3	1	J	3	1	2	K	3	2	2
△計	27	20	10		21	22	17	승計	19	23	18	合計	27	21	18

2. Plasmid construction





Expression plasmid map

Conclusion and discussion

No significant differences were noted in the results neither among the 4 kinds of organic acid composition nor between the two different kinds of concentration of the same organic acid.

But we can make sure about the unique taste-modifying activity of the MIR with 4 kinds of the organic acid composition.

We constructed the plasmid pLTSSP6HisM to express the recombination miraculin in yeast cells.

We'll purify the MIR and re-MIR, and do the CD spectroscopic analysis to try to know the conformational change of miraculin acid-response. At last we'll try to figure out the 3D protein structure of miraculin.