

ふりがな 氏名	いしい じゅん 石井 純	職名	教授
部局 講座 教育研究分野名	科学技術イノベーション研究科 バイオ・環境講座 バイオプロダクション分野	学位称号	博士（工学）
略歴（学歴，職歴，教育歴，所属機関，学会及び社会における活動，受賞）			
年 月	（学 歴）		
1998年3月	私立西大和学園高等学校 卒業		
2002年3月	神戸大学工学部応用化学科 卒業		
2004年3月	神戸大学大学院自然科学研究科応用化学専攻 博士前期課程 修了		
2008年3月	神戸大学大学院自然科学研究科分子物質科学専攻 博士後期課程 修了		
年 月	（職 歴）		
2008年4月	神戸大学自然科学系先端融合研究環重点研究部 学術推進研究員		
2008年8月	神戸大学自然科学系先端融合研究環重点研究部 特命助教		
2012年4月	神戸大学自然科学系先端融合研究環重点研究部 特命准教授		
2013年4月	神戸大学自然科学系先端融合研究環重点研究部 准教授		
2016年4月	神戸大学大学院科学技術イノベーション研究科 准教授		
2018年7月	神戸大学先端バイオ工学研究センター 准教授		
2018年7月	（神戸大学大学院科学技術イノベーション研究科 兼任）		
2024年10月	（神戸大学工学部応用化学科 兼任）		
年 月	（受 賞）		
2021年10月	第73回日本生物工学会大会トピックス賞（日本生物工学会）		
2022年10月	令和4年度学長表彰・財務貢献者（神戸大学）		
2023年6月	令和5年度研究科優秀教育賞（神戸大学大学院科学技術イノベーション研究科）		
2023年10月	令和5年度学長表彰・財務貢献者（神戸大学）		
2024年9月	第47回生物工学奨励賞・照井賞（日本生物工学会）		
2024年10月	令和6年度学長表彰・財務貢献者（神戸大学）		

教 育 研 究 上 の 業 績

(著 書)

1. 戸谷吉博, 松田史生, 石井純, 原清敬, 微生物を活用した有用物質の製造技術, 第1章 バイオものづくりに資する微生物と育種技術, 第4節 「光をエネルギー源として利用する有用物質生産大腸菌の開発」シーエムシー出版, 2023年5月31日
2. 柘植謙爾, 石井純, 近藤昭彦, ゲノム編集技術を応用した製品開発とその実用化 ～研究開発動向・課題解決策・技術予測と市場展望～, 第4章 ゲノム編集によるスマートセルインダストリーの技術開発とその課題解決, 第7節 「長鎖DNA合成技術による有用物質生産微生物の構築とその課題解決」技術情報協会, 2021年2月26日
3. 石井純, 西晶子, 他6名, スマートセルインダストリー -微生物細胞を用いた物質生産の展望- 第1編 ハイスループット合成・分析・評価技術, 第2章 ハイスループット微生物構築・評価技術, 第1節 「微生物を用いた物質生産とハイスループット微生物構築技術」シーエムシー出版, 2018年6月
4. Nakamura Y, Kondo A, Ishii J*. (2018) 「Peripheral Membrane Proteins (担当: Chapter 3 “Biosensing techniques in yeast: G-protein signaling and protein-protein interaction assays for monitoring ligand stimulation and oligomer formation of heterologous GPCRs”)」 InTechOpen Book Chapter
5. 石井純, 近藤昭彦, 化学便覧 応用化学編 第7版 第VII編 バイオ科学技術, 第26章 バイオマテリアル, 第3節 「ヒドロキシ化合物」丸善出版, 2014年1月
6. 石井純, 荻野千秋, 近藤昭彦, 次世代医薬開発に向けた抗体工学の最前線, 第II編 抗体の改変技術, 第4章 親和性の向上, 第4節 「酵母による抗体フラグメントおよび抗体様結合性タンパク質の改変技術」シーエムシー出版, 2012年12月
7. 石井純, 田中勉, 他2名, シングルセル解析の最前線 第3章 細胞内生体分子群の実測定量解析, 第4節 「フローサイトメトリーと GFP レポーターによる G 蛋白質シグナルのシングルセル解析」シーエムシー出版, 2010年3月
8. 石井純, 近藤昭彦, 一細胞定量解析の最前線 -ライフサーベイヤ構築に向けて- 第3章 細胞内生体分子群の動態シグナルの解析, 第6節 「酵母細胞内シグナル定量解析の創薬への応用」シーエムシー出版, 2006年12月

(学 術 論 文)

(a. 学会誌, 専門誌等に掲載された論文)

1. Tominaga M, Shima Y, Nozaki K, Ito Y, Someda M, Shoya Y, Hashii N, Obata C, Matsumoto-Kitano M, Suematsu K, Matsukawa T, Hosoya K, Hashiba N, Kondo A, Ishii J*. (2024) Designing strong inducible synthetic promoters in yeasts. *Nat. Commun.* 15(1): 10653
2. Asama R, Tominaga M, Ito S, Ito Y, Takemura K, Sakuraba S, Katsurada K, Fukuda N, Kondo A, Ishii J*. (2024) Screening of protein-based inhibitors for the intracellular domain of epidermal growth factor receptor by directed evolution using the yeast Gy recruitment system. *J. Biosci. Bioeng.* 138(5): 375-381
3. Asama R, Liu CJS, Tominaga M, Cheng YR, Nakamura Y, Kondo A, Wang HY*, Ishii J*. (2024) Droplet-based microfluidic platform for detecting agonistic peptides that are self-secreted by yeast expressing a G-protein-coupled receptor. *Microb. Cell Fact.* 23(1): 104
4. Matsuyama C, Seike T, Okahashi N, Niide T, Hara KY, Hirono-Hara Y, Ishii J, 他3名. (2023) Metabolome analysis of metabolic burden in *Escherichia coli* caused by overexpression of green fluorescent protein and delta-rhodopsin. *J. Biosci. Bioeng.* 137(3): 187-194
5. Morita K, Takenaka M, Tomita K, Ishii J, 他8名. (2023) Nanoscopic lignin mapping on cellulose nanofibers via scanning transmission electron microscopy and atomic force microscopy. *Cellulose* 30: 11357-11367

6. Sugimura M, Seike T, Okahashi N, Izumi Y, Bamba T, **Ishii J**, Matsuda F. (2023) Improved 2,3-butanediol production rate of metabolically engineered *Saccharomyces cerevisiae* by deletion of *RIM15* and activation of pyruvate consumption pathway. *Int. J. Mol. Sci.* 24(22): 16378
7. Yatabe F, Seike T, Okahashi N, **Ishii J**, Matsuda F. (2023) Improvement of ethanol and 2,3-butanediol production in *Saccharomyces cerevisiae* by ATP wasting. *Microb. Cell Fact.* 22(1): 204
8. Arai T*, Wada M, Nishiguchi H, Takimura Y, **Ishii J***. (2023) Inducer-free recombinant protein production in *Trichoderma reesei*: Secretory production of endogenous enzymes and heterologous nanobodies using glucose as the sole carbon source. *Microb. Cell Fact.* 22(1): 103
9. Takekana M, Yoshida T, Yoshida E, Ono S, Horie S, Vavricka CJ, Hiratani M, Tsuge K, **Ishii J**, 他 3 名. (2023) Online SFE-SFC-MS/MS colony screening: A high-throughput approach for optimizing (-)-limonene production. *J. Chromatogr. B* 1215: 123588
10. Sano M, Tanaka R, Kamata K, Hirono-Hara Y, **Ishii J**, 他 4 名. (2022) Conversion of mevalonate to isoprenol using light energy in *Escherichia coli* without consuming sugars for ATP supply. *ACS Synth. Biol.* 11(12): 3966-3972
11. Morita K, Nishimura Y, **Ishii J**, Maruyama T. (2022) Micelle-like nanoassemblies of short peptides create antimicrobial selectivity in a conventional antifungal drug. *ACS Appl. Nano Mater.* 6(2): 1432-1440
12. Otsuka K, Seike T, Toya Y, **Ishii J**, 他 3 名. (2022) Evolutionary approach for improved proton pumping activity of heterologous rhodopsin expressed in *Escherichia coli*. *J. Biosci. Bioeng.* 134(6): 484-490
13. Ito Y, Ishigami M, Hashiba N, Nakamura Y, Terai G, Hasunuma T, **Ishii J***, Kondo A*. (2022) Avoiding entry into intracellular protein degradation pathways by signal mutations increases protein secretion in *Pichia pastoris*. *Microb. Biotechnol.* 15(9): 2364-2378
14. Toya Y, Hirono-Hara Y, Hirayama H, Kamata K, Tanaka R, Sano M, Kitamura S, Otsuka K, Abe-Yoshizumi R, Tsunoda SP, Kikukawa H, Kandori H, Shimizu H, Matsuda F, **Ishii J**, Hara KY. (2022) Optogenetic reprogramming of carbon metabolism using light-powering microbial proton pump systems. *Metab. Eng.* 72: 227-236
15. Isogai S, Tominaga M, Kondo A, **Ishii J***. (2022) Plant flavonoid production in bacteria and yeasts. *Front. Chem. Eng.* 4: 880694 [Review]
16. Tominaga M, Miyazaki K, Hataya S, Mitsui Y, Kuroda S, Kondo A, **Ishii J***. (2022) Enhanced squalene production by modulation of pathways consuming squalene and its precursor. *J. Biosci. Bioeng.* 134(1): 1-6
17. Ito Y, Ishigami M, Terai G, Nakamura Y, Hashiba N, Nishi T, Nakazawa H, Hasunuma T, Asai K, Umetsu M, **Ishii J***, Kondo A*. (2022) A streamlined strain engineering workflow with genome-wide screening detects enhanced protein secretion in *Komagataella phaffii*. *Commun. Biol.* 5(1): 561
18. Tominaga M, Kondo A, **Ishii J***. (2022) Engineering of synthetic transcriptional switches in yeast. *Life* 12(4): 557 [Review]
19. Vavricka CJ, Takahashi S, Watanabe N, Takenaka M, Matsuda M, Yoshida T, Suzuki R, Kiyota H, Li J, Minami H, **Ishii J**, 他 4 名. (2022) Machine learning discovery of missing links that mediate alternative branches to plant alkaloids. *Nat. Commun.* 13(1): 1405
20. Morita K, Seike T, **Ishii J**, 他 2 名. (2022) Improvement of 2,3-butanediol production by dCas9 gene expression system in *Saccharomyces cerevisiae*. *J. Biosci. Bioeng.* 133(3): 208-212
21. Nishi T, Ito Y, Nakamura Y, Yamaji T, Hashiba N, Tamai M, Yasohara Y, **Ishii J***, Kondo A*. (2022) One-step *in vivo* assembly of multiple DNA fragments and genomic integration in *Komagataella phaffii*. *ACS Synth. Biol.* 11(2): 644-654
22. Katsurada K, Tominaga M, Kaishima M, Kato H, Matsuno T, Ogino C, Kondo A, **Ishii J***, Takayama K. (2021) Constitutive cell surface expression of ZZ domain for the easy preparation of yeast-based immunosorbents. *J. Gen. Appl. Microbiol.* 67(6): 265-268

23. Yoshida E*, Kojima M, Suzuki M, Matsuda F*, Shimbo K, Onuki A, Nishio Y, Usuda Y, Kondo A, **Ishii J***. (2021) Increased carvone production in *Escherichia coli* by balancing limonene conversion enzyme expression via targeted quantification concatamer proteome analysis. *Sci. Rep.* 11(1): 22126
24. Isogai S, Okahashi N, Asama R, Nakamura T, Hasunuma T, Matsuda F, **Ishii J***, Kondo A*. (2021) Synthetic production of prenylated naringenins in yeast using promiscuous microbial prenyltransferases. *Metab. Eng. Commun.* 12: e00169.
25. Shoji S, Yamaji T, Makino H, **Ishii J**, Kondo A. (2021) Metabolic design for selective production of nicotinamide mononucleotide from glucose and nicotinamide. *Metab. Eng.* 167-177
26. Tominaga M, Nozaki K, Umeno D, **Ishii J***, Kondo A. (2021) Robust and flexible platform for directed evolution of yeast genetic switches. *Nat. Commun.* 12(1): 1846
27. Nakamura Y, Asama R, Tabata T, Morita K, Maruyama T, Kondo A, **Ishii J***. (2021) Comparative analyses of site-directed mutagenesis of human melatonin MTNR1A and MTNR1B receptors using a yeast fluorescent biosensor. *Biotechnol. Bioeng.* 118(2): 863-876
28. Ito Y, Terai G, Ishigami M, Hashiba N, Nakamura Y, Bamba T, Kumokita R, Hasunuma T, Asai K, **Ishii J***, Kondo A*. (2020) Exchange of endogenous and heterogeneous yeast terminators in *Pichia pastoris* to tune mRNA stability and gene expression. *Nucleic Acids Res.* 48(22): 13000-13012
29. Nishimura Y, Ezawa R, Morita K, Nakayama M, **Ishii J**, 他 3 名. (2020) *In vivo* evaluation of the Z_{HER2}-BNC/LP carrier encapsulating an anticancer drug and a radiosensitizer. *ACS Appl. Bio Mater.* 3(11): 7743-7751
30. Morita K, Matsuda F, Okamoto K, **Ishii J**, 他 2 名. (2019) Repression of mitochondrial metabolism for cytosolic pyruvate-derived chemical production in *Saccharomyces cerevisiae*. *Microb. Cell Fact.* 18(1): 177
31. Vavricka CJ, Yoshida T, Kuriya Y, Takahashi S, Ogawa T, Ono F, Agari K, Kiyota H, Li J, **Ishii J**, 他 5 名. (2019) Mechanism-based tuning of insect 3,4-dihydroxyphenylacetaldehyde synthase for synthetic bioproduction of benzylisoquinoline alkaloids. *Nat. Commun.* 10(1): 2015
32. Tamano K, Cox RS 3rd, Tsuge K, Miura A, Itoh A, **Ishii J**, 他 3 名. (2019) Heterologous production of free dihomog- γ -linolenic acid by *Aspergillus oryzae* and its extracellular release via surfactant supplementation. *J. Biosci. Bioeng.* 127(4): 451-457
33. Itoh H, Matsui M, Miyamura Y, Takeda I, **Ishii J**, 他 4 名. (2018) Biosynthesis of novel statins by combining heterologous genes from *Xylaria* and *Aspergillus*. *ACS Synth. Biol.* 7(12): 2783-2789
34. Ito Y, Watanabe T, Aikawa S, Nishi T, Nishiyama T, Nakamura Y, Hasunuma T, Okubo Y, **Ishii J***, Kondo A*. (2018) Deletion of DNA ligase IV homolog confers higher gene targeting efficiency on homologous recombination in *Komagataella phaffii*. *FEMS Yeast Res.* 18(7): foy074
35. Restu WK, Nishida Y, Yamamoto S, **Ishii J**, Maruyama T. (2018) Short oligopeptides for biocompatible and biodegradable supramolecular hydrogels. *Langmuir* 34(27): 8065-8074
36. Nakamura Y, Nishi T, Noguchi R, Ito Y, Watanabe T, Nishiyama T, Aikawa S, Hasunuma T, **Ishii J***, Okubo Y, Kondo A*. (2018) A stable, autonomously replicating plasmid vector containing *Pichia pastoris* centromeric DNA. *Appl. Environ. Microbiol.* 84(15): e02882-17
37. **Ishii J***, Morita K, 他 6 名, Matsuda F*. (2018) A pyruvate carbon flux tugging strategy for increasing 2,3-butanediol production and reducing ethanol subgeneration in the yeast *Saccharomyces cerevisiae*. *Biotechnol. Biofuels* 11: 180
38. Hashi H, Nakamura Y, **Ishii J***, Kondo A. (2018) Modifying expression modes of human neurotensin receptor type 1 alters sensing capabilities for agonists in yeast signaling biosensor. *Biotechnol. J.* 13(4): e1700522
39. Nishimura Y, Matsui T, **Ishii J**, Kondo A. (2018) Metabolic engineering of the 2-ketobutyrate biosynthetic pathway for 1-propanol production in *Saccharomyces cerevisiae*. *Microb. Cell Fact.* 17(1): 38

40. Nambu-Nishida Y, Sakihama Y, **Ishii J**, 他 2 名. (2018) Selection of yeast *Saccharomyces cerevisiae* promoters available for xylose cultivation and fermentation. *J. Biosci. Bioeng.* 125(1): 76-86
41. Prima A, Hara KY, Djohan AC, Kashiwagi N, Kahar P, **Ishii J**, 他 6 名. (2017) Glutathione production from mannan-based bioresource by mannanase/mannosidase expressing *Saccharomyces cerevisiae*. *Bioresour. Technol.* 245(Pt B): 1400-1406
42. Hara KY, Kobayashi J, Yamada R, Sasaki D, Kuriya Y, Hirono-Hara Y, **Ishii J**, 他 2 名. (2017) Transporter engineering in biomass utilization by yeast. *FEMS Yeast Res.* 17(7): fox061 (Mini review)
43. Morita K, Nomura Y, **Ishii J**, 他 3 名. (2017) Heterologous expression of bacterial phosphoenol pyruvate carboxylase and Entner-Doudoroff pathway in *Saccharomyces cerevisiae* for improvement of isobutanol production. *J. Biosci. Bioeng.* 124(3): 263-270
44. Fukuda N, Kaishima M, **Ishii J**, Honda S. (2017) Positive detection of GPCR antagonists using a system for inverted expression of a fluorescent reporter gene. *ACS Synth. Biol.* 6(8): 1554-1562
45. Fukutani Y, **Ishii J**, 他 4 名. (2017) Split luciferase complementation assay for the analysis of G protein-coupled receptor ligand response in *Saccharomyces cerevisiae*. *Biotechnol. Bioeng.* 114(6): 1354-1361
46. Ryo S, **Ishii J**, 他 5 名. (2017) Positive feedback genetic circuit incorporating a constitutively active mutant Gal3 into yeast GAL induction system. *ACS Synth. Biol.* 6(6): 928-935
47. Nishimura Y, Ezawa R, **Ishii J**, 他 2 名. (2017) Affibody-displaying bio-nanocapsules effective in EGFR, typical biomarker, expressed in various cancer cells. *Bioorg. Med. Chem. Lett.* 27(2): 336-341
48. Fukuda N, Kaishima M, **Ishii J**, 他 2 名. (2016) Continuous crossbreeding of sake yeasts using growth selection systems for α -type and α -type cells. *AMB Express* 6(1): 45
49. Inokuma K, Bamba T, **Ishii J**, 他 3 名. (2016) Enhanced cell-surface display and secretory production of cellulolytic enzymes with *Saccharomyces cerevisiae* Sed1 signal peptide. *Biotechnol. Bioeng.* 113(11): 2358-2366
50. Kaishima M, **Ishii J***, 他 3 名. (2016) Expression of varied GFPs in *Saccharomyces cerevisiae*: codon optimization yields stronger than expected expression and fluorescence intensity. *Sci. Rep.* 6: 35932
51. Nakamura Y, Hashimoto T, **Ishii J***, Kondo A. (2016) Dual-color reporter switching system to discern dimer formations of G-protein-coupled receptors using Cre/*loxP* site-specific recombination in yeast. *Biotechnol. Bioeng.* 113(10): 2178-2190
52. **Ishii J**, Okazaki F, Djohan AC, 他 11 名. (2016) From mannan to bioethanol: cell surface co-display of β -mannanase and β -mannosidase on yeast *Saccharomyces cerevisiae*. *Biotechnol. Biofuels* 9(1): 188
53. Yamamoto K, Hara KY, Morita T, Nishimura A, Sasaki D, **Ishii J**, 他 3 名. (2016) Enhancement of astaxanthin production in *Xanthophyllomyces dendrorhous* by efficient method for the complete deletion of genes. *Microb. Cell Fact.* 15(1): 155
54. Shen L, Nishimura Y, Matsuda F, **Ishii J**, Kondo A. (2016) Overexpressing enzymes of the Ehrlich pathway and deleting genes of the competing pathway in *Saccharomyces cerevisiae* for increasing 2-phenylethanol production from glucose. *J. Biosci. Bioeng.* 122(1): 34-39
55. Nakamura Y, **Ishii J**, Kondo A. (2016) Current techniques for studying oligomer formations of G-protein-coupled receptors using mammalian and yeast cells. *Curr. Med. Chem.* 23(16): 1638-1656
[Review]
56. Hattan J, Shindo K, Ito T, Shibuya Y, Watanabe A, Tagaki C, Ohno F, Sasaki T, **Ishii J**, 他 2 名. (2016) Identification of a novel hedyeryol synthase gene isolated from *Camellia brevistyla* flowers and floral scent of *Camellia* cultivars. *Planta* 243(4): 959-972
57. Tsuge Y, Kudou M, Kawaguchi H, **Ishii J**, 他 2 名. (2016) FudC, a protein primarily responsible for furfural detoxification in *Corynebacterium glutamicum*. *Appl. Microbiol. Biotechnol.* 100(6): 2685-2692
58. Hasunuma T, **Ishii J**, Kondo A. (2015) Rational design and evolutionary fine tuning of *Saccharomyces cerevisiae* for biomass breakdown. *Curr. Opin. Chem. Biol.* 1-9 [Review]

59. Hiraoka R, Funasaki Y, **Ishii J**, Maruyama T. (2015) Rational design of degradable polyanion for layer-by-layer assembly for encapsulation and release of cationic functional biomolecules. *Chem. Commun.* 51(98): 17447-17450
60. Kaishima M, **Ishii J***, 他 2 名. (2015) Gy recruitment systems specifically select PPI and affinity-enhanced candidate proteins that interact with membrane protein targets. *Sci. Rep.* 5: 16723
61. Nakamura Y, **Ishii J**, Kondo A. (2015) Applications of yeast-based signaling sensor for characterization of antagonist and analysis of site-directed mutants of the human serotonin 1A receptor. *Biotechnol. Bioeng.* 112(9): 1906-1915
62. Ida K, **Ishii J**, 他 3 名. (2015) Eliminating the isoleucine biosynthetic pathway to reduce competitive carbon outflow during isobutanol production by *Saccharomyces cerevisiae*. *Microb. Cell Fact.* 14: 62
63. Inokuma K, **Ishii J**, 他 4 名. (2015) Complete genome sequence of *Kluyveromyces marxianus* NBRC1777, a nonconventional thermotolerant yeast. *Genome Announc.* 3(2). pii: e00389-15
64. Takenaka M, Miyachi Y, **Ishii J**, 他 2 名. (2015) The mapping of yeast's G-protein coupled receptor with an atomic force microscope. *Nanoscale* 7(11): 4956-4963
65. Suzuki H, **Ishii J**, 他 2 名. (2015) Polyamino acid display on cell surfaces enhances salt and alcohol tolerance of *Escherichia coli*. *Biotechnol. Lett.* 37(2): 429-435
66. Bin Mohamed Suffian IF, Nishimura Y, Morita K, Nakamura-Tsuruta S, Al-Jamal KT, **Ishii J**, 他 2 名. (2015) Mutation of arginine residues to avoid non-specific cellular uptakes for hepatitis B virus core particles. *J. Nanobiotechnol.* 13(1): 15
67. Fukutani Y, Hori A, Tsukada S, Sato R, **Ishii J**, 他 3 名. (2015) Improving the odorant sensitivity of olfactory receptor-expressing yeast with accessory proteins. *Anal. Biochem.* 471: 1-8
68. Inokuma K, Yoshida T, **Ishii J**, 他 2 名. (2015) Efficient co-displaying and artificial ratio control of α -amylase and glucoamylase on the yeast cell surface by using combinations of different anchoring domains. *Appl. Microbiol. Biotechnol.* 99(4): 1655-1663
69. Nakamura Y, **Ishii J**, Kondo A. (2014) Construction of a yeast-based signaling biosensor for human angiotensin II type 1 receptor via functional coupling between Asn295-mutated receptor and Gpa1/G_{i3} chimeric G α . *Biotechnol. Bioeng.* 111(11): 2220-2228
70. Tsuge Y, Hori Y, Kudou M, **Ishii J**, 他 2 名. (2014) Detoxification of furfural in *Corynebacterium glutamicum* under aerobic and anaerobic conditions. *Appl. Microbiol. Biotechnol.* 98(20): 8675-8683
71. Nishimura Y, **Ishii J**, 他 2 名. (2014) Genetic engineering of bio-nanoparticles for drug delivery: A Review. *J Biomed Nanotechnol* 10(9): 2063-2085 [Review]
72. Kaishima M, Fukuda N, **Ishii J**, Kondo A. (2014) Desired alteration of protein affinities: competitive selection of protein variants using yeast signal transduction machinery. *PLoS One* 9(9): e108229
73. Nakamura Y, **Ishii J**, Kondo A. (2014) Signaling assays for detection of human G-protein-coupled receptors in yeast. *Bio Protoc* 4(16) <http://www.bio-protocol.org/e1206>
74. **Ishii J**, Kondo T, 他 4 名. (2014) Three gene expression vector sets for concurrently expressing multiple genes in *Saccharomyces cerevisiae*. *FEMS Yeast Res.* 14(3): 399-411
75. Nishimura Y, Takeda K, Ezawa R, **Ishii J**, 他 2 名. (2014) A display of pH-sensitive fusogenic GALA peptide facilitates endosomal escape from a Bio-nanocapsule via an endocytic uptake pathway. *J. Nanobiotechnology* 12: 11
76. Nakamura Y, Takemoto N, **Ishii J**, Kondo A. (2014) Simultaneous method for analyzing dimerization and signaling of G-protein-coupled receptor in yeast by dual-color reporter system. *Biotechnol. Bioeng.* 111(3): 586-596
77. **Ishii J**, Oda A, 他 5 名. (2014) Microbial fluorescence sensing for human neurotensin receptor type 1 using G α -engineered yeast cells. *Anal. Biochem.* 446: 37-43
78. Matsuda F, **Ishii J**, 他 4 名. (2013) Increased isobutanol production in *Saccharomyces cerevisiae* by eliminating competing pathways and resolving cofactor imbalance. *Microb. Cell Fact.* 12: 119

79. Nakamura Y, **Ishii J**, Kondo A. (2013) Bright fluorescence monitoring system utilizing *Zoanthus* sp. green fluorescent protein (*ZsGreen*) for human G-protein-coupled receptor signaling in microbial yeast cells. *PLoS One* 8(12): e82237
80. Matsuda F, Shirai T, **Ishii J**, Kondo A. (2013) Regulation of central carbon metabolism in *Saccharomyces cerevisiae* by metabolic inhibitors. *J. Biosci. Bioeng.* 116(1): 59-64
81. Nishimura Y, Mieda H, **Ishii J**, 他 3 名. (2013) Targeting cancer cell-specific RNA interference by siRNA delivery using a complex carrier of affibody-displaying bio-nanocapsules and liposomes. *J. Nanobiotechnology* 11: 19
82. Nakamura Y, **Ishii J**, Kondo A. (2013) Rapid, facile detection of heterodimer partners for target human G-protein-coupled receptors using a modified split-ubiquitin membrane yeast two-hybrid system. *PLoS One* 8(6): e66793
83. Nishimura Y, Takeda K, **Ishii J**, 他 2 名. (2013) An affinity chromatography method used to purify His-tag-displaying bio-nanocapsules. *J. Virol. Methods* 189(2): 393-396
84. Hasunuma T, Okazaki F, Okai N, Hara KY, **Ishii J**, Kondo A. (2013) A review of enzymes and microbes for lignocellulosic biorefinery and the possibility of their application to consolidated bioprocessing technology. *Bioresour. Technol.* 135: 513-522 [Review]
85. **Ishii J**, Yoshimura K, 他 2 名. (2013) Reduction of furan derivatives by overexpressing NADH-dependent Adh1 improves ethanol fermentation using xylose as sole carbon source with *Saccharomyces cerevisiae* harboring XR-XDH pathway. *Appl. Microbiol. Biotechnol.* 97(6): 2597-2607
86. Nishimura Y, Mimura W, Suffian IFM, Amino T, **Ishii J**, 他 2 名. (2013) Granting specificity for breast cancer cells using a hepatitis B core particle with a HER2-targeted affibody molecule. *J. Biochem.* 153(3): 251-256
87. Suga H, Matsuda F, Hasunuma T, **Ishii J**, Kondo A. (2013) Implementation of a transhydrogenase-like shunt to counter redox imbalance during xylose fermentation in *Saccharomyces cerevisiae*. *Appl. Microbiol. Biotechnol.* 97(4): 1669-1678
88. Kondo A, **Ishii J**, 他 3 名. (2013) Development of microbial cell factories for bio-refinery through synthetic bioengineering. *J. Biotechnol.* 163(2): 204-216 [Review]
89. Nishimura Y, **Ishii J**, 他 3 名. (2012) Complex carriers of affibody-displaying bio-nanocapsules and composition-varied liposomes for HER2-expressing breast cancer cell-specific protein delivery. *J. Drug Target.* 20(10): 897-905
90. Fukutani Y, **Ishii J**, 他 3 名. (2012) An improved bioluminescence-based signaling assay for odor sensing with a yeast expressing a chimeric olfactory receptor. *Biotechnol. Bioeng.* 109(12): 3143-3151
91. Matsuda F, Kondo T, Ida K, Tezuka H, **Ishii J**, Kondo A. (2012) Construction of an artificial pathway for isobutanol biosynthesis in the cytosol of *Saccharomyces cerevisiae*. *Biosci. Biotechnol. Biochem.* 76(11): 2139-2141
92. **Ishii J**, Moriguchi M, 他 4 名. (2012) Improved identification of agonist-mediated G α_i -specific human G-protein-coupled receptor signaling in yeast cells by flow cytometry. *Anal. Biochem.* 426(2): 129-133
93. Ryo S, **Ishii J**, 他 3 名. (2012) Transplantation of the GAL regulon into G-protein signaling circuitry in yeast. *Anal. Biochem.* 424(1): 27-31
94. Kondo T, Tezuka H, **Ishii J**, 他 3 名. (2012) Genetic engineering to enhance the Ehrlich pathway and alter carbon flux for increased isobutanol production from glucose by *Saccharomyces cerevisiae*. *J. Biotechnol.* 159(1-2): 32-37
95. **Ishii J**, Yoshimoto N, 他 5 名. (2012) Cell wall trapping of autocrine peptides for human G-protein-coupled receptors on the yeast cell surface. *PLoS One* 7(5): e37136
96. Hara KY, Kim S, Kiriya K, Yoshida H, Arai S, **Ishii J**, 他 3 名. (2012) An energy-saving glutathione production method from low-temperature cooked rice using amylase-expressing *Saccharomyces cerevisiae*. *Biotechnol J* 7(5): 686-689

97. Fukutani Y, Nakamura T, Yorozu M, **Ishii J**, 他 2 名. (2012) The N-terminal replacement of an olfactory receptor for the development of a yeast-based biomimetic odor sensor. *Biotechnol. Bioeng.* 109(1): 205-212
98. Nishimura Y, Shishido T, **Ishii J**, 他 3 名. (2012) Protein-encapsulated bio-nanocapsules production with ER membrane localization sequences. *J. Biotechnol.* 157(1): 124-129
99. Fukuda N, **Ishii J**, 他 2 名. (2011) Amplification of agonist stimulation of human G-protein-coupled receptor signaling in yeast. *Anal. Biochem.* 417(2): 182-187
100. Fukuda N, **Ishii J**, Kondo A. (2011) G γ recruitment system incorporating a novel signal amplification circuit to screen transient protein-protein interactions. *FEBS J.* 278(17): 3086-3094
101. Matsuda F, Furusawa C, Kondo T, **Ishii J**, 他 2 名. (2011) Engineering strategy of yeast metabolism for higher alcohol production. *Microb. Cell Fact.* 10: 70
102. Hasunuma T, Sanda T, Yamada R, Yoshimura K, **Ishii J**, Kondo A. (2011) Metabolic pathway engineering based on metabolomics confers acetic and formic acid tolerance to a recombinant xylose-fermenting strain of *Saccharomyces cerevisiae*. *Microb. Cell Fact.* 10(1): 2
103. Tanino T, Hotta A, Ito T, **Ishii J**, 他 6 名. (2010) Construction of a xylose-metabolizing yeast by genome integration of xylose isomerase gene and investigation of the effect of xylitol on fermentation. *Appl. Microbiol. Biotechnol.* 88(5): 1215-1221
104. Togawa S, **Ishii J**, 他 4 名. (2010) Importance of asparagine residues at positions 13 and 26 on the amino-terminal domain of human somatostatin receptor subtype-5 in signalling. *J. Biochem.* 147(6): 867-873
105. Iguchi Y, **Ishii J**, 他 6 名. (2010) Control of signalling properties of human somatostatin receptor subtype-5 by additional signal sequences on its amino-terminus in yeast. *J. Biochem.* 147(6): 875-884
106. **Ishii J**, Fukuda N, 他 3 名. (2010) Protein-protein interactions and selection: yeast-based approaches that exploit guanine nucleotide-binding protein signaling. *FEBS J.* 277(9): 1982-1995. [Minireview] (Selected as cover)
107. Fukuda N, **Ishii J**, 他 3 名. (2010) The competitor-introduced G γ recruitment system, a new approach for screening affinity-enhanced proteins. *FEBS J.* 277(7): 1704-1712
108. Tanaka T, Masunari S, **Ishii J**, 他 4 名. (2010) Displaying non-natural, functional molecules on yeast surfaces via biotin-streptavidin interaction. *J. Biotechnol.* 145(1): 79-83
109. **Ishii J**, Izawa K, 他 7 名. (2009) A simple and immediate method for simultaneously evaluating expression level and plasmid maintenance in yeast. *J. Biochem.* 145(6): 701-708
110. Fukuda N, **Ishii J**, 他 3 名. (2009) Construction of a novel detection system for protein-protein interactions using yeast G-protein signaling. *FEBS J.* 276(9): 2636-2644
111. Shibasaki S, Sakata K, **Ishii J**, 他 2 名. (2008) Development of a yeast protein fragment complementation assay (PCA) system using dihydrofolate reductase (DHFR) with specific additives. *Appl. Microbiol. Biotechnol.* 80(4): 735-743
112. **Ishii J**, Tanaka T, 他 6 名. (2008) Yeast-based fluorescence reporter assay of G protein-coupled receptor signaling for flow cytometric screening: *FAR1*-disruption recovers loss of episomal plasmid caused by signaling in yeast. *J. Biochem.* 143(5): 667-674
113. Fukuda N, **Ishii J**, 他 4 名. (2008) Rapid and efficient selection of yeast displaying a cell-surface protein using magnetic nanoparticles. *Biotechnol. Prog.* 24(2): 352-357
114. Fukuda N, **Ishii J**, 他 4 名. (2007) High-efficiency recovery of target cells using improved yeast display system for detection of protein-protein interactions. *Appl. Microbiol. Biotechnol.* 76(1): 151-158
115. Shibasaki S, Kawabata A, **Ishii J**, 他 6 名. (2007) Construction of a novel synergistic system for production and recovery of secreted recombinant proteins by the cell surface engineering. *Appl. Microbiol. Biotechnol.* 75(4): 821-828

116. **Ishii J**, Matsumura S, 他 5 名. (2006) Quantitative and dynamic analyses of G protein-coupled receptor signaling in yeast using Fus1, enhanced green fluorescence protein (EGFP), and His3 fusion protein. *Biotechnol. Prog.* 22(4): 954-960

* : 責任著者

(学 術 報 告 等)

1. 富永将大, 近藤昭彦, **石井純** 「酵母における遺伝子スイッチ進化工学ワークフローの「ロボスタ化」」 生物物理, 第 64 巻, 第 3 号, 144-146 (2024 年 6 月号)
2. **石井純**, 新型コロナで変わる時代の実験自動化・遠隔化 「微生物での発酵生産と実験の自動化」 実験医学 (羊土社), 2021 年 1 月号, 第 39 巻, 第 1 号, 8-12 (2020 年 12 月)
3. **石井純**, 荒木通啓, 他 5 名 「合成生物学によるモノづくり微生物のデザインに向けて」 生物工学会誌, 第 93 巻, 第 9 号, 523-526 (2015 年 9 月)
4. **石井純**, 蓮沼誠久, 他 2 名 「バイオリファイナリー社会に向けた燃料・化学品生産」 安全工学 (安全工学会 会誌), 第 52 巻, 第 4 号, 249-255 (2013 年 8 月)
5. **石井純**, 蓮沼誠久, 他 2 名 「革新的なモノづくり実現のための「合成生物学」」 生物工学会誌, 第 91 巻, 第 6 号, 314-318 (2013 年 6 月)

(上記以外に 12 編)

(学 術 講 演)

1. **Ishii J**. Yeast Genetic Parts and Tools for Synthetic Biology and Cell Factories. *2025 Asian Synthetic Biology Association (ASBA) Meeting (ASBA2025)*. 2025, Jan 6-10, Hotel Fort Canning, Singapore 【招待講演】
2. **石井純** 「微生物代謝の理解と最適化に向けたバイオ DX 基盤の構築」 第 75 回日本生物工学会大会 シンポジウム (生命科学研究の革新に向けたバイオ DX の挑戦), 2023 年 9 月 3-5 日, 名古屋大学東山キャンパス 【招待講演】
3. **石井純** 「酵母における合成生物学基盤の開発と物質生産への展開」 日本農芸化学会 2023 年度大会 シンポジウム (微生物によるものづくり研究の最前線 ~宿主機能開発から社会実装に向けた取り組みまで), 2023 年 3 月 14-17 日, オンライン開催 【招待講演】
4. **石井純** 「バイオエコノミー社会実現に向けた合成生物学の基盤開発と微生物での物質生産への展開」 SONY “Life Science Spring Webinar 2022”, 2022 年 3 月 23 日, オンライン開催 【招待講演】
5. **Ishii J**, Tabata T, Nakamura T, Kato H, Asama R, Nakamura Y, Kondo A. Construction of metabolite sensor using yeast signal transduction machinery for monitoring melatonin production. *The 10th Symposium on Innovative BioProduction Taichung (iBioT2019)*. 2019, Nov 7-9, Tunghai University, Taichung, Taiwan 【招待講演】

(上記以外に 294 編)