

| No | Client | Client Lab | JRPP Member | Collaborative Research Item |
|----|-------------------------------------|--|-----------------|---|
| 1 | Clarence Yeung | Molecular Ligand Target Research Team | Akiko Idei | Screening compound management and microdispensing |
| 2 | Nicole Case | Molecular Ligand Target Research Team | Akiko Idei | Screening compound management and microdispensing |
| 3 | Pham Lien | Molecular Ligand Target Research Team | Akiko Idei | Screening compound management and microdispensing |
| 4 | Ryoung Shin | Environmental Response Research Unit | Makoto Kawatani | Preparation of chemical beads |
| 5 | Ken Matsumoto | Chemical Genomics Research Group | Makoto Kawatani | Preparation of chemical beads |
| 6 | Makoto Muroi | Biomolecular Characterization Unit | Makoto Kawatani | Preparation of chemical beads |
| 7 | Yoko Yashiroda | Molecular Ligand Target Research Team | Makoto Kawatani | Preparation of chemical beads |
| 8 | Kentaro Nishiyama | Molecular Bioregulation Research Team | Norio Kudo | Crystal structure determination |
| 9 | Hamish Craig | Biomacromolecules Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 10 | Hamish Craig | Biomacromolecules Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 11 | Jagat Chhipi Shrestha | Chemical Genomics Research Group | Naoshi Dohmae | Quantitative proteomic analysis |
| 12 | Pham, Lien Thi Kim | Molecular Ligand Target Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 13 | Schneider-Poetsch, Tilman Hans Jorg | Chemical Genomics Research Group | Naoshi Dohmae | Quantitative proteomic analysis |
| 14 | Shamitha Rao YAGI | Biomacromolecules Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 15 | Xin Tong | Cell Function Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 16 | Shunji Takahashi | Natural Product Biosynthesis Research Unit | Naoshi Dohmae | Quantitative proteomic analysis |
| 17 | Ken Matsumoto | Chemical Genomics Research Group | Naoshi Dohmae | Quantitative proteomic analysis |
| 18 | Minoru Ueda | Plant Genomic Network Research Team | Naoshi Dohmae | Amino acid analysis_protein absolute quantification / Quantitative proteomic analysis |
| 19 | Yoko Chiba | Biofunctional Catalyst Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 20 | Yoko Chiba | Biofunctional Catalyst Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 21 | Kentaro Nakaminami | Plant Genomic Network Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 22 | Kentaro Nakaminami | Plant Genomic Network Research Team | Naoshi Dohmae | Quantitative proteomic analysis |
| 23 | Yu Zheng | Natural Product Biosynthesis Research Unit | Naoshi Dohmae | Quantitative proteomic analysis |
| 24 | Keisuke Fujiyama | Natural Product Biosynthesis Research Unit | Naoshi Dohmae | Quantitative proteomic analysis |
| 25 | Hiromi Suzuki | Dormancy and Adaptation Research Unit | Naoshi Dohmae | Measurement of compounds and macromolecules by MALDI-TOF MS |
| 26 | Ken Matsumoto | Chemical Genomics Research Group | Hiroyuki Hirano | Providing compounds |
| 27 | Fumiyuki Shirai | Drug Discovery Chemistry Platform Unit | Hiroyuki Hirano | Providing compounds |
| 28 | Hiroo Koyama | Drug Discovery Chemistry Platform Unit | Hiroyuki Hirano | Providing compounds |
| 29 | Akiko Idei | Drug Discovery Seed Compounds Exploratory Unit | Hiroyuki Hirano | Providing compounds |
| 30 | Akiko Idei | Drug Discovery Seed Compounds Exploratory Unit | Hiroyuki Hirano | Providing compounds |
| 31 | Akiko Idei | Drug Discovery Seed Compounds Exploratory Unit | Hiroyuki Hirano | Providing compounds |
| 32 | Pham Lien | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |
| 33 | Pham Lien | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |
| 34 | Yoko Yashiroda | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |

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| 35 | Yoko Yashiroda | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |
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| 37 | Yoko Yashiroda | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |
| 38 | Yoko Yashiroda | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |
| 39 | Yoko Yashiroda | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |
| 40 | Mie Kubota | Molecular Ligand Target Research Team | Hiroyuki Hirano | Providing compounds |
| 41 | Akiko Idei | Drug Discovery Seed Compounds Exploratory Unit | Hiroyuki Hirano | Providing compounds |
| 42 | Akiko Idei | Drug Discovery Seed Compounds Exploratory Unit | Hiroyuki Hirano | Providing compounds |
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| 50 | Hiroo Koyama | Drug Discovery Chemistry Platform Unit | Hiroyuki Hirano | Providing compounds |
| 51 | Yu Zheng | Natural Product Biosynthesis Research Unit | Hiroyuki Hirano | Providing compounds |
| 52 | Kosuke Dodo | Catalysis and Integrated Research Group | Ken Matsumoto | Validation after CRISPR screening |
| 53 | Yoichi Yamada | Green Nanocatalysis Research Team | Atsuya Muranaka | Measurement and analysis of ultraviolet-visible-near-infrared, infrared absorption, and fluorescence spectra |
| 54 | Yoichi Yamada | Green Nanocatalysis Research Team | Atsuya Muranaka | Molecular structure analysis by quantum chemical calculation |
| 55 | Masayoshi Nishiura | Advanced Catalysis Research Group | Atsuya Muranaka (Takashi Nakamura) | Measurement and analysis of solid-state NMR |
| 56 | Takayuki Motoyama | Plant Immunity Research Group | Atsuya Muranaka (Toshihiko Nogawa) | Qualitative analysis by mass spectrometry / NMR measurement and analysis |
| 57 | Yoko Chiba | Biofunctional Catalyst Research Team | Atsuya Muranaka (Hiroyuki Koshino) | NMR measurement and analysis |
| 58 | Katsuyuki Sakai | Natural Product Biosynthesis Research Unit | Atsuya Muranaka (Hiroyuki Koshino, Naoya Takahashi) | NMR measurement and analysis / Molecular structure analysis by quantum chemical calculation / Synthesis of organic compounds, absolute configuration determination by asymmetric total synthesis |
| 59 | Ju Yeon Moon | Environmental Response Research Unit | Makoto Muroi | Protein identification |
| 60 | Ju Yeon Moon | Environmental Response Research Unit | Makoto Muroi | Protein identification |
| 61 | Tilman Schneider-Poetsch | Chemical Genomics Research Group | Makoto Muroi | Proteomic analysis |

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| 62 | Katsuhiko Sekimata | Drug Discovery Chemistry Platform Unit | Seiji Matsuoka | Binding affinity measurement of biomolecular interactions by microscale thermophoresis (MST) |
| 63 | Takayoshi Awakawa | Chemical Biology and Biosynthesis Research Team | Seiji Matsuoka | Kinetic analysis of biomolecular interactions by surface plasmon resonance (SPR) / Protein stability evaluation by differential scanning fluorescence spectroscopy (nanoDSF) |
| 64 | Naoshi Dohmae | Biomolecular Characterization Unit | Atsuya Muranaka (Naoya Takahashi) | Absolute configuration determination by asymmetric total synthesis |
| 65 | Akihiro Ito | Chemical Genomics Research Group | Naoshi Dohmae | Quantitative proteomic analysis |
| 66 | Shintaro Kawamura | Catalysis and Integrated Research Group | Yoko Yashiroda | High-throughput screening (Yeast chemical genomics analysis) |
| 67 | Yoshihiro Sohtome | Catalysis and Integrated Research Group | Yoko Yashiroda | High-throughput screening (Yeast chemical genomics analysis) |
| 68 | Ikko Takahashi | Advanced Organic Synthesis Research Team | Yoko Yashiroda | High-throughput screening (Yeast chemical genomics analysis) |