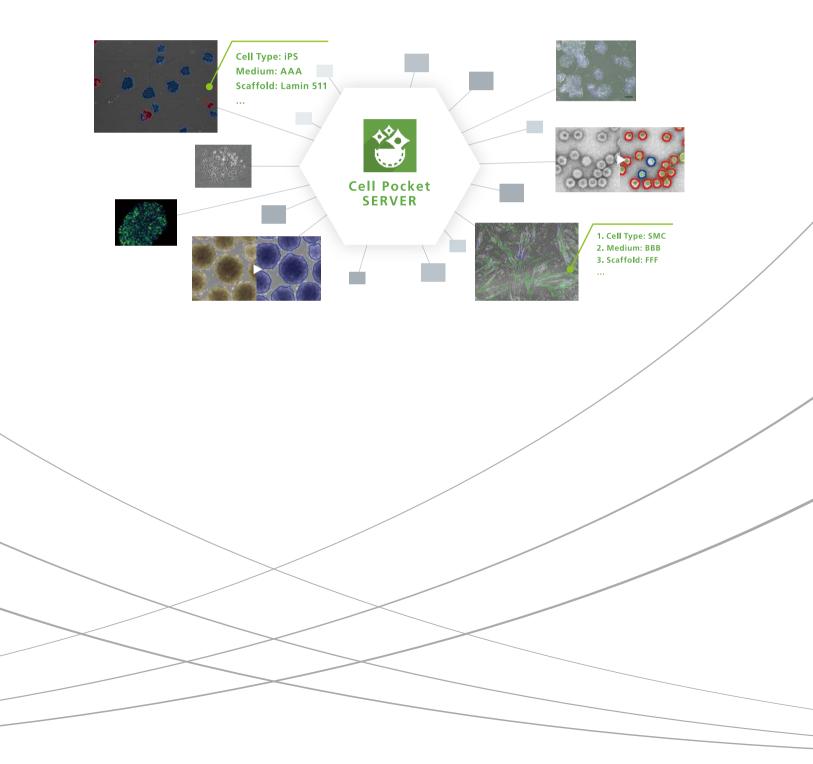


Web Application Supporting Cell Observation

Cell Pocket

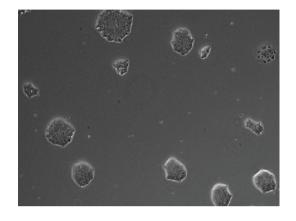


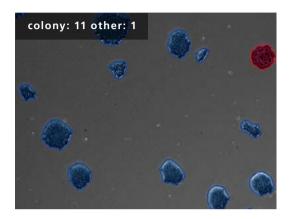
New Experience in Cell Image Analysis

— Using AI to display image information objectively and quantitatively —

Example

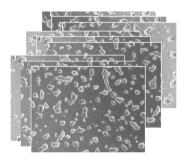
Detection of abnormal colonies



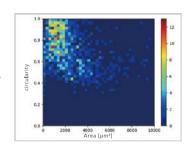


Example

Quantification and graphing of the morphological features of cell colonies









Quantify the visual state of cells from ———— P. 4 various aspects

- Deep learning technology can offer the quantification of cell status.
- Easy quantifying and graphing with customizable analysis recipes.



Centralized management of all data from ——— P. cell observation work

- Easy to gather various cell culture information such as culture conditions, image data, and analyzed data.
- This is a Web application for up to 20 users.
- Centralized management facilitates sharing data and knowledge among members.



User interface specialized for cell culture management

P. 7

POINT 3

- Cell image data can be easily linked to culture conditions.
- The user interface for culture management is simple and easy to understand and supports a user's cell analysis work.





Process flow for image analysis using Al

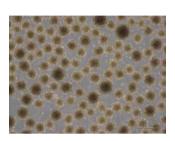


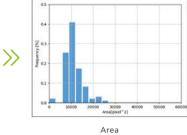


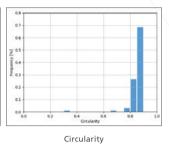
Set data analysis goals.

Clearly identify the desired analysis results.

Example: Evaluating the variability of spheroid area and circularity



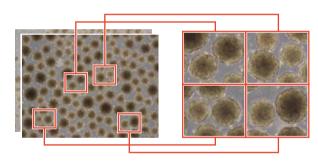


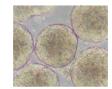


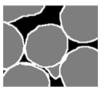
02

Acquire AI training data.

Acquire images necessary for obtaining desired results. Verification is possible from a minimum of about 10 image data sets.







Target image

Labeled image

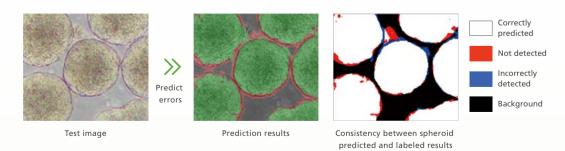
Note: Manually labeled or color-coded images are used as labeled images.



Train AI model and assess performance.

Train the AI model and assess its performance with test images. Cell Pocket™ automatically assigns a test image, numerically evaluates the entire test image, and indicates any regions of the test image where errors were predicted.

The figures below are used to assess whether spheroid regions were correctly predicted.



Note: Processes for calculating area and circularity values from predicted spheroid regions are specified in data analysis recipes indicated on the next page.

In this example, the results confirm whether spheroid regions can be correctly identified.

Quantify the visual state of cells from various aspects

Analysis recipes can be made in three steps

Constructed recipes can be used routinely for the same purpose



Preparation of trained model*

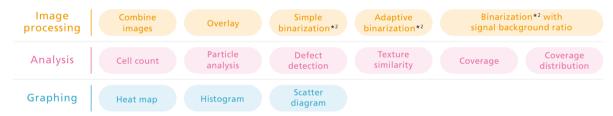
Cell Pocket equips the preparation function of trained models to deduce cell or colony regions by using a user's image data.





Selection of analysis processing scripts

Cell Pocket contains 14 kinds of commonly used analysis processing scripts. By assembling them, a variety of analysis recipes can be constructed. It is also possible to describe and add scripts on your own.

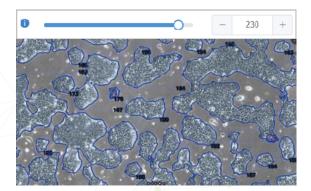


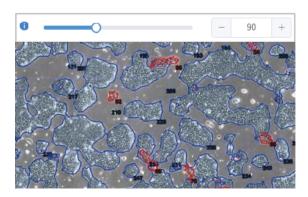
^{*2 &}quot;Binarization" means image processing for a discriminating area of interest or that of no interest. For example, this is used for discriminating nucleus and no cell region.



Optimization of parameters for analysis processing scripts

Parameters for analysis processing scripts can be adjusted with the user interface. Figures below show analysis examples before and after changing the parameters for the threshold of defect cells.





Analysis examples

Analysis of colony morphology distribution

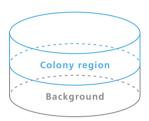
Trained model

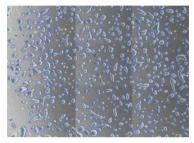
ombine images

Coverage

Particle analysis

After combining nine images, colony regions are discriminated from the background, and then the diameter, area and circularity of each colony and coverage can be quantitated.





#	Diameter	Area	•••	Aspect ratio
1	16.1	107.8		5.4
2	37.3	223.1		10.2
3	57.1	420.3		8.9
		:		
242	32.7	180.2		6.8
243	44.2	267.9		6.2

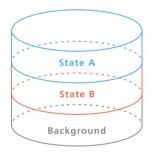
Coverage: 29 %

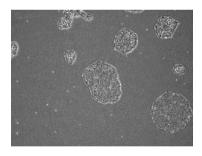
Analysis of texture similarity

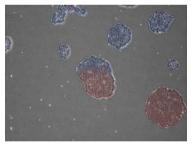
Trained model

Texture similarity

After training, a slight difference in pattern information on the cell texture can be detected and quantified.







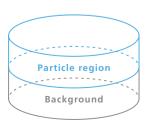
Similarity score with state A: 53 %

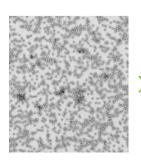
Particle classification by morphological analysis and characteristic quantities

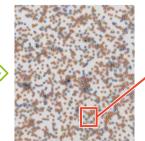
Trained model

Particle analysis

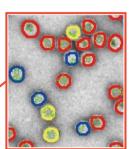
By classifying particle types based on characteristic quantities, such as the shape, color, or brightness of particles, particles with particular characteristics can be counted.







>>>



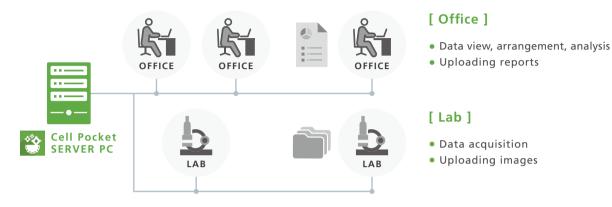


Centralized management of all cell culture data in your lab

Manage data via a Web application

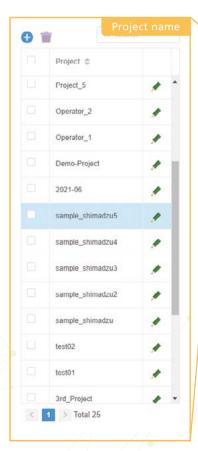
After installing a server PC in your office, you can use this system via a web browser.

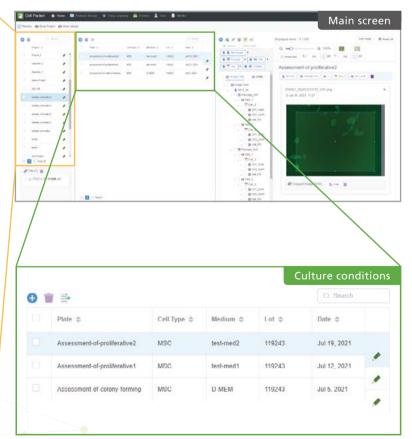
Note: Up to 20 users per one server PC



At-a-glance viewer for cell culture information

- Displays culture conditions, acquired data, and analyzed results at a glance
- Reduces labor time for comparing with past results, searching acquired data, and sharing the experimental progress



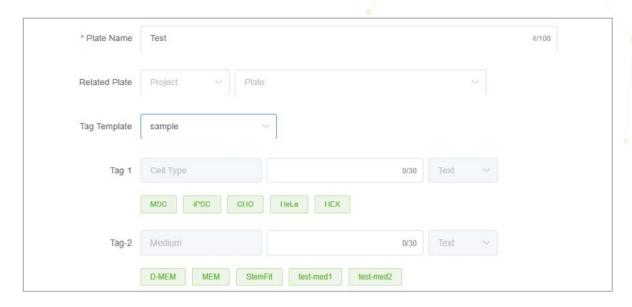




User interface specialized for cell culture management

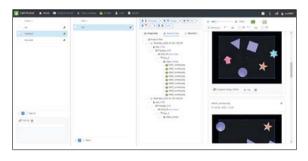
Input culture conditions with ease and without omission

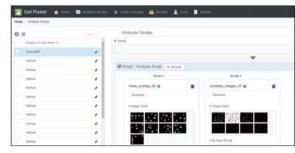
- Input by selecting specific cell culture conditions, such as cell line and culture medium.
- Cell culture information input from lab members can be consistent using the "template Input" function.
- New templates can be created and administrative users can edit the template.

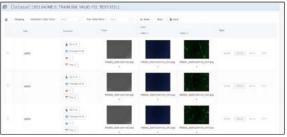


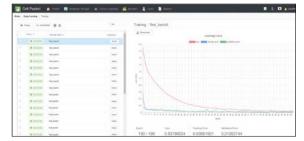
Support data management for utilizing AI

- Data traceability is ensured by keeping a record of data sets, training histories, data analysis recipes, and data analysis results.
- It effortlessly achieves data management, which is even more important due to the use of Al.









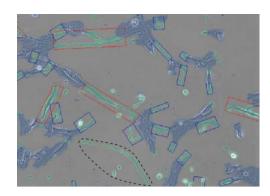
Related Software

Morphology Analysis Plug-in for Mesenchymal Stem Cells

This application extracts the filamentous pseudopodia from cellular regions, quantifies the long axis length and its ratio in comparison to the total, and outputs a histogram.

This product is an analysis script developed through collaborative research with Dr. Shin Kawamata at the Research & Development Center for Cell Therapy, Foundation for Biomedical Research and Innovation in Kobe, Japan.

Refer to the flyer C297-E140 for details.



Counted as long pseudopodia Counted as short pseudopodia

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