



(51) International Patent Classification:

G16B 25/10 (2019.01) G06N 3/02 (2006.01)
G16B 40/30 (2019.01)

(21) International Application Number:

PCT/SG2024/050008

(22) International Filing Date:

04 January 2024 (04.01.2024)

(25) Filing Language:

English

(26) Publication Language:

English

(30) Priority Data:

10202300148U 18 January 2023 (18.01.2023) SG

(71) Applicant: **AGENCY FOR SCIENCE, TECHNOLOGY AND RESEARCH** [SG/SG]; 1 Fusionopolis Way, #20-10, Connexis North Tower, Singapore 138632 (SG).

(72) Inventors: **LONG, Yahui**; c/o Genome Institute of Singapore, 60 Biopolis Street, #02-01, Singapore 138672 (SG).
CHEN, Jinmiao; c/o Genome Institute of Singapore, 60 Biopolis Street, #02-01, Singapore 138672 (SG).

(74) Agent: **DAVIES COLLISON CAVE ASIA PTE. LTD.**; 10 Collyer Quay #07-01, Ocean Financial Centre, Singapore 049315 (SG).

(81) Designated States (unless otherwise indicated, for every kind of national protection available): AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BN, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CV, CZ, DE, DJ, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IQ, IR, IS, IT, JM, JO, JP, KE, KG, KH, KN, KP, KR, KW, KZ, LA, LC, LK, LR, LS, LU, LY, MA, MD, MG, MK, MN, MU, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PA, PE, PG, PH, PL, PT, QA, RO, RS, RU, RW, SA, SC, SD, SE, SG, SK, SL, ST, SV, SY, TH, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, WS, ZA, ZM, ZW.

(84) Designated States (unless otherwise indicated, for every kind of regional protection available): ARIPO (BW, CV, GH, GM, KE, LR, LS, MW, MZ, NA, RW, SC, SD, SL, ST, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, RU, TJ, TM), European (AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, ME, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, KM, ML, MR, NE, SN, TD, TG).

Declarations under Rule 4.17:

— of inventorship (Rule 4.17(iv))

(54) Title: SPATIALLY INFORMED CLUSTERING, INTEGRATION, AND DECONVOLUTION OF SPATIAL TRANSCRIPTOMICS WITH GRAPHST

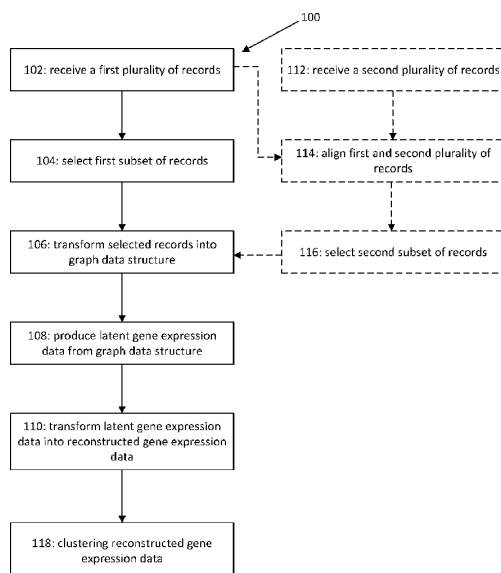


Figure 1

(57) Abstract: Systems and methods for processing spatial transcriptomics data and generating insights relating cell and tissue states. The system and methods including spatial clustering, integration of data from multiple tissue samples and integration of scRNA-sequence data and spatial transcriptomics data. The systems and methods incorporate graph self-supervised contrastive learning and graph neural networks to perform spatial transcriptomics data processing.

Published:

- *with international search report (Art. 21(3))*
- *before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments (Rule 48.2(h))*