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## In vitro cancer cell line classification using pattern recognition approach based on metabolite profiling (Article)

Jeffree, A.I.<sup>a</sup> [✉](#), Omar, M.I.<sup>a</sup>, Has-Yun Hashim, Y.<sup>b</sup>, Zakaria, A.<sup>a</sup>, Thriumani, R.<sup>a</sup>, Shakaff, A.Y.M.<sup>a</sup> [👤](#)<sup>a</sup>Centre of Excellence for Advanced Sensor Technology (CEASTech), Universiti Malaysia Perlis, Perlis, Malaysia<sup>b</sup>Cell and Tissue Culture Engineering Lab, Department of Biotechnology Engineering, International Islamic University Malaysia, Gombak, Selangor, Malaysia

## Abstract

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This study aims to evaluate the feasibility of metabolite profiling for the characterisation and discrimination volatile compounds using the pattern recognition from in vitro cancer cell lines, which are lung, breast and colon cancer together with the blank medium as a control group. This study implemented the A549 (lung), MCF7 (breast) and HCT116 (colon). Cells were harvested and maintained until they grow as monolayer adherent and reach confluence 70-90% before sampling. The volatiles profile from the targeted cell line was established using headspace solid phase microextraction coupled to gas chromatography-mass spectrometry (HS-SPME/GCMS). Multivariate data analysis employed principal component analysis (PCA) to better visualise the subtle similarities and the differences among these data sets. A total of 116 volatile organic compounds were detected focused on a limited range of retention time from 3rd until 17th minutes, and 33 compounds were recognized as targeted compounds (peak area>1%). According to both results, the score and the loading plot explained 83% of the total variance. The volatiles compound has shown to be significantly distinguished among cancerous and control group based on metabolite profiling using pattern recognition approach. © 2018 Universiti Teknikal Malaysia Melaka. All rights reserved.

## Reaxys Database Information

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## References (32)

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- 1 Dyzmann-Sroka, A., Malicki, J.  
Cancer incidence and mortality in the Greater Poland Region-Analysis of the year 2010 and future trends (2014) *Reports Pract. Oncol. Radiother.*  
Jul.

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J.  
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occupational medicine practiceTaeger, D. , Gawrych, K. ,  
Brüning, T.  
(2016) *International Journal of  
Occupational Medicine and  
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- 2 Ferlay, J., Soerjomataram, I., Dikshit, R., Eser, S., Mathers, C., Rebelo, M., Parkin, D.M., (...), Bray, F.  
Cancer incidence and mortality worldwide: Sources, methods and major patterns in GLOBOCAN 2012

(2015) *International Journal of Cancer*, 136 (5), pp. E359-E386. Cited 7518 times.  
[http://onlinelibrary.wiley.com/journal/10.1002/\(ISSN\)1097-0215](http://onlinelibrary.wiley.com/journal/10.1002/(ISSN)1097-0215)  
doi: 10.1002/ijc.29210

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---

- 3 Visvader, J.E.  
Cells of origin in cancer

(2011) *Nature*, 469 (7330), pp. 314-322. Cited 684 times.  
doi: 10.1038/nature09781

[View at Publisher](#)

---

- 4 Li, H., Kim, B.-R., Wu, R.  
Identification of quantitative trait nucleotides that regulate cancer growth: A simulation approach

(2006) *Journal of Theoretical Biology*, 242 (2), pp. 426-439. Cited 5 times.  
doi: 10.1016/j.jtbi.2006.03.010

[View at Publisher](#)

---

- 5 Zhang, A., Sun, H., Xu, H., Qiu, S., Wang, X.  
Cell metabolomics

(2013) *OMICS A Journal of Integrative Biology*, 17 (10), pp. 495-501. Cited 34 times.  
doi: 10.1089/omi.2012.0090

[View at Publisher](#)

---

- 6 Gowda, G.A.N., Zhang, S., Gu, H., Asiago, V., Shanaiah, N., Raftery, D.  
Metabolomics-based methods for early disease diagnostics

(2008) *Expert Review of Molecular Diagnostics*, 8 (5), pp. 617-633. Cited 268 times.  
<http://www.expert-reviews.com/doi/pdf/10.1586/14737159.8.5.617>  
doi: 10.1586/14737159.8.5.617

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---

- 7 Halama, A., Guerrouahen, B.S., Pasquier, J., Diboun, I., Karoly, E.D., Suhre, K., Rafii, A.  
Metabolic signatures differentiate ovarian from colon cancer cell lines

(2015) *Journal of Translational Medicine*, 13 (1), art. no. 223. Cited 16 times.  
<http://www.translational-medicine.com/home/>  
doi: 10.1186/s12967-015-0576-z

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---

- 8 Shulaev, V.  
Metabolomics technology and bioinformatics

(2006) *Briefings in Bioinformatics*, 7 (2), pp. 128-139. Cited 200 times.  
doi: 10.1093/bib/bbl012

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